Greater Paraburdoo Iron Ore Hub Proposal (Assessment No. 2189)

CMS17501 Greater Paraburdoo Iron Ore Hub – Terrestrial Fauna comments

Table 1: Department of Water and Environmental Regulation Comments on Terrestrial Fauna

Item	Торіс	DWER Comment	Rio Tinto Response
Verteb	rate Fauna		
1	Surveys	The surveys for vertebrate fauna are adequate to predict the impacts of the proposal.	Noted.
2	Environmental Management Plan (Appendix 4)	 Issues relating to the management of impacts to Ghost bat (<i>Macroderma gigas</i>): For caves that are within 300 metres (m) of proposed operations (i.e., caves 6, 16, 17 and 18) a 100 m mining restriction zone (MRZ) is proposed (MRZ [Table 1.2]). However, the term 'mining exclusion zone' (MEZ) is used for the Pilbara leaf-nosed bat (PLNB) roost at the Ratty Springs cave structure which is also larger at 250 m (Table 1.2). The EMP includes a management-based target of "ensure no significant long-term decline in ghost bat usage of high value habitat in the Development Envelope, attributable to the Proposal" (Table 1). Acoustic monitoring of 10 significant roost sites at Western range and four regional roosts will be undertaken to "indicate how Ghost bats use caves within the Greater Paraburdoo region (e.g., diurnal versus maternal/ caves at western range vs other regional caves)" that will be analysed bi-annually and species tracking of individuals annually for five to ten nights (Table 2-4). However, it is unclear if: recording acoustic data alone will be 	Noted. The Proponent has opted to apply a 250 m Mining Exclusion Zone (M at Ratty Spring. No direct disturbance from activities associated with to the restricted shape and size of the Ratty Springs roost, accessing also not possible to collect temperature or humidity data from the ro Spring roost has been taken as a precautionary approach due to the data from inside the roost chamber cannot be collected. A threshold conservative vibration value) will also be set for the Ratty Springs roo For Ghost Bat roosts at Western Range, the Proponent has applied 1 no direct clearing for mining activities (e.g. no drilling) will be pern potentially extend into the MRZ. The 100m MRZ for Ghost Bat roosts is of suitable caves within the Development Envelope (18 caves, the Development Envelope (18 caves, the
		 able to differentiate between roost usage e.g., diurnal vs maternal; the regional roosts are intended to be reference (control) sites; and whether the proposed biannual analysis of monitoring data will be adequate to measure significant changes to cave use by ghost bats. It appears that mining activities will overlap the proposed baseline establishment period. The baseline commenced in 2020/2021 and will proceed to 2022/2023 and mining at Western Ranges is proposed to commence in 2023 (p.17). 	Proposal), the documented highly transitory nature of Ghost Bats at V of Ghost Bats at Western Range, and the low resident Ghost bat popu It is important to note however that the environmental objective of structural integrity of the caves is not impacted by the Proposal so tha enduring habitat for PLNB and Ghost bat beyond the life of the Propo The MEZ and MRZ's have also been modified since they were present
		 Issues relating to the management of impacts to Pilbara leaf-nosed bat (<i>Rhinonicteris aurantius</i> [Pilbara form]): A MEZ of 250 m is proposed around the Pilbara leaf-nosed bat (PLNB) roost at Ratty Springs (Table 1.2). However, the term 'MRZ' is used for the retained ghost bat cave 	around the known cave dimensions rather than around the cave en roost. The MEZ over the PLNB Ratty Spring roost has also been incr approach based on significance of the roost site and acknowledging install/collect temperature or humidity data from the roost chamber.
		 structures. The Response to Submissions (RTS) states that no vibration monitoring and no vibration limits will be assigned for the Ratty Springs PLNB roost as the site does not share the same geology with the proposed mine area (Response to Submissions, Part A.4). Instead, annual visual inspections of the external structure are proposed (Table 1.2). Annual inspections may not be adequate as they may not detect changes to the internal stability of the cave structure within a reasonable period to implement actions. In addition, no evidence has been provided to support 	Responses to issues relating to the management of impacts to Ghost. The Proponent notes that generally, acoustic recording only prov numbers. However, from baseline monitoring since 2019, accurate usage and the low number of Ghost Bats residing in the Western Rang monitoring. The results of acoustic monitoring have been verified a hormone analysis and visual cave inspections.
		 the statement that the differences in geology between the Ratty Springs cave structure and the proposed mine area will limit the transmission of vibrations and there will be no impact to the PLNB colony. The PLNB colony at the Ratty Springs cave will be monitored using acoustic data. Acoustic surveys (call counts) will provide an indication of the activity level of the PLNB colony. However, this technique alone does not represent an accurate estimate of colony size (abundance) and behaviour as PLNB habit is to fly repeatedly in and out of the roost structure after dusk, causing an increase in 	Continuous acoustic monitoring data is available from 2019 for the 10 2021a). As construction activities are set to commence in 2023, this p for this species at Western Range. Baseline monitoring will continue u the Proponent will ensure the baseline data collection period does no Proposal. Maternity cave classification via acoustic monitoring has been accur roosting in caves during the maternity period. The results of this meth and visual cave inspections conducted within Western Range and

MEZ) over the Pilbara Leaf-nosed Bat (PLNB) roost th the Proposal are permitted within a MEZ. Due the roost chamber is not possible; it is therefore roost chamber. The larger MEZ around the Ratty biological importance of the site and noting that I of 10mm/s peak particle velocity (PPV) (the most pst.

100m Mining Restriction Zones (MRZ). In MRZ's, mitted, however some indirect disturbance may is considered sufficient based on the high number majority of which will not be impacted by the Western Range, no evidence of roost site fidelity ulation (approximately 10 resident individuals).

f the MEZ and MRZ's is the same. To ensure the at these important features will remain, providing osal.

ted in the ERD. The zones now represent a buffer ntrance, providing protection around the entire reased from 150 m to 250 m as a precautionary g accessing the roost chamber is not possible to

Bat

vides relative abundance rather than individual estimates of Ghost Bat numbers including cave ges (n=10), have been confirmed through acoustic against other monitoring methods including scat

LO Ghost Bat caves specified in the EMP (Rio Tinto provides a minimum of four years of baseline data until implementation of the Proposal commences; not overlap with the implementation phase of the

arately inferred from the presence of continued nod are consistent with the scat hormone analysis d across other Rio Tinto Ghost Bat monitoring

Item	Торіс	DWER Comment	Rio Tinto Response
		activity that is not a true reflection of bat abundance (Cramer et al., 2016; Bullen	programs currently being undertaken in concert with Robert Bullen (Ba
		and Reiffer, 2020). Acoustic data in combination with infrared or thermal video	for the species than other methods that require cave entry Radio Fre
		censusing techniques should be used to approximate trends in colony size from the	analysis, especially for a species such as the Ghost Bat that is easily dist
		call data (Bullen and Reiffer, 2020).	
		• An environmental outcome of "no significant long-term decline in the Pilbara	Currently there are few Ghost Bat roosts known and monitored at a
		Leaf-nosed bat population utilising the permanent maternity roost at Ratty Springs,	20-50km), to act as direct control/reference sites. Identifying such locat
		attributable to the Proposal" has been assigned for the PLNB colony at the Ratty	as Panhandle, Turee and Western Turner Syncline. Data from the region
		Springs cave (Table 1). The trigger and threshold criteria are based on a 'lower call	monitoring undertaken at several other Rio Tinto sites across the Pilbara
		limit' (LCL [Summary Table 1]) that is derived from the long-term average of nightly	population and seasonal trends, with the regional roosts considered
		call counts (Figure A.2) and is based on several other PLNB maternity roosts. It is	commencement of work to identify Ghost Bat roosts at a regional scale
		unclear whether this is adequate to detect changes in PLNB activity and colony size,	
		within and between years. In addition, setting criteria limits based on lower limits	Recording of Ghost Bats acoustic data in the 10 Ghost Bat caves sp
		will only capture change in minimum values. Therefore, an increase or decrease in	continuous throughout the year. Analysis of data will be undertaken or
		the number of peak events (e.g. breeding period July to March) or a seasonal shift	activities) or biannual (all other retained Ghost bat roosts) basis. This n
		may not be captured using the proposed criteria.	up outside of the cave with microphones situated in the roost char
		• Acoustic data will be analysed on a quarterly basis, therefore there is a risk that	recording echolocation and social calls of this species, without the impar
		trigger and threshold criteria may be exceeded prior to detection and management	not show roost site fidelity, specifically in the Western Ranges area, cont
		actions being implemented (Table 2.6).	to monitor the presence of Gnost Bats and to measure changes in cav
		It appears that mining activities will overlap the proposed baseline establishment	continual data collection, it allows for changes to be identified and attrict
		period. The baseline commenced in 2020/2021 and will proceed to 2022/2023 and	(unlike other monitoring methods such as scat analysis which only allo
		mining at Western Ranges is proposed to commence in 2023 (p.17).	over the collection period). It is also noted that the low number of Gho
			cave usage throughout the year. The Proponent does not consider that
		Actions required:	notential changes considering these factors
		• The definitions for the MRZ and the MEZ should include the activities that will be (used to be activities to be activitities to be activitities to be a	potential changes considering these factors.
		restricted (or excluded) and those that will be unrestricted (or included). In	Responses to issues relating to the management of impacts to Pilhara k
		the MPZ and MEZ (i.e. 100 m for the rate and short bet as a structures of 250 m	A geotechnical assessment of the Ratty Springs PLNB roost will be cond
		for the Datty Spring DLND cave structure)	roost's geotechnical sensitivity to structural instability. As a precaution
		The MDZ and MEZ chould take into consideration the full extent of the cave	most conservative vibration value) will be set for the Ratty Springs roo
			biological importance. A vibration monitor will be located at the roost o
		 The assumption that the Patty Springs cave structure will not be subject to vibration. 	monitoring equipment) and vibration monitoring requirements and trigg
		• The assumption that the Katty springs cave structure will not be subject to vibration	(Table 2.3 and Table 2.4, Rio Tinto 2021a) and captured in the site's blas
		site should be verified using blast testing monitoring and modelling. In addition	
		as the Ratty Springs cave structure is a maternal roost site for the PINB it is	Initial blast modelling using Australian Standards (k = 1140 and b -1.6)
		recommended that vibration limits are set based on blast modelling and testing and	<10mm/s PPV is achievable using a mix of support and production
		are incorporated in the Environmental Management Plan (EMP) using outcome-	Brockman Iron geology data from the Proponent's other operations in the
		based management provisions.	be achieved (k = 731 and b -1.28 as the 95% confidence level). Both initial
		• It is recommended that the Proponent should consult with the Department of	the nearest mine pits at approximately 350m; these pits have now bee
		Biodiversity, Conservation and Attractions (DBCA) regarding a minimum baseline	proposed pits at Western Range to the Ratty Springs PLNB roost now g
		for the ghost bat and PLNB. However, it is anticipated that a minimum baseline of	initial models are presented in Figure 1 and Figure 2 below. Detail
		two years (or two breeding seasons) should be established for the ghost bat and	undertaken prior to any pit blasting associated with Western Range bei
		PLNB colonies, prior to ground disturbing activities. The Proponent should	
		demonstrate the adequacy of any proposed baseline.	
		• Provide information on how the activity data (call counts) for PLNB will be verified	
		to estimate the size of the colony e.g. using infrared camera analysis (or other	
		methods).	
		• The monitoring should be designed using Before-After-Control-Impact analysis and	
		the EMP should include the statistical and analytical methods (EPA 2020) to	
		measure significant changes in the PLNB and ghost bat colonies.	

BatCall WA). This method is also far less invasive requency Identification (RFID) tagging and scat sturbed.

a regional scale from Western Range (within ations will be required, specifically in areas such onal roosts (once located) in addition to acoustic ara will allow for a bioregion wide comparison of d reference sites. The Proponent is targeting e in 2022.

specified in the EMP (Rio Tinto 2021a) will be on a quarterly (Ghost bat roosts within 300 m of method uses permanent acoustic recorders set amber, allowing for an increased likelihood of act of cave entry by personnel. As Ghost Bats do ntinuous acoustic monitoring is the best method aves usage across the site. As this method uses ibuted to activities associated with the Proposal low identification of whether a site was utilised nost Bats residing in the Western Ranges (n=10), tern Ranges leads to high variability in individual at more frequent data analysis will help identify

leaf-nosed bat

ducted prior to mining activities to establish the onary measure, a threshold of 10mm/s PPV (the post for the duration of the Proposal due to its or close by (due to space constraints with other ggers/thresholds have been included in the EMP ast management controls for the roost.

as the 95% confidence level has indicated that n drill and blast parameters. Modelling using the Pilbara also indicates that <10mm/s PPV can nitial models assumed 14W-16W and 20W were een removed from the design, with the nearest greater than 2,800m away. The results of both niled modelling using site specific data will be eing undertaken.

Item	Торіс	DWER Comment	Rio	Tinto Response	
		Clarify if the regional roosts monitored for ghost bats (i.e. Pan Handle, Turee Creek			
		and Karijini) are intended to be reference sites.	-	Million Bar Hadam for	
		• Revise the trigger and threshold criteria based on LCL to detect changes in the	ĩ	vibration Prediction for	Peak Part
		seasonal use of the Ratty Springs roost by PLNB (i.e. Table 2.6). In addition, consider	H		Scaled Di
		including an upper call limit (and relevant trigger and threshold criteria) and	nto		
		increasing the frequency of the call analysis during critical periods (such as during	0		
		the breeding cycle for the PLNB - July to March).		10	100
		• Consider including scat analysis (for DNA and hormones) as a monitoring technique		K PPV value when SD = 1 731 mm/s 10 b Local rock exponent -1.28 (-1.6 typ) 10	
		for PLNB, as has been used for the West Angelas Project (ghost bat).			
		 For consistency with other projects, the following terms (related to bats) should be 		MIC Max charge in 8ms 150 kg D Distance from blast 350.6 m	
		defined (e.g. TSSC 2016).		D Distance non blast 200.0 m	
		Roost: is the area within a roost structure (i.e. cave, adit or artificial		SD Scaled Distance 28.6 m/vkg	100
		structure) where the bats actually aggregate during roosting.		PPV Peak particle velocity 10.0 mm/s	
		 Colony: group of individuals occupying a roost structure at a particular. 		(s,m	
		time.		Dist (m) SD (m/vkg) PPV (mm/s) E	-
		 Population: all individuals of the species in a defined geographical area. 		10 0.8 947.6	
		For example, for the PINB this is usually the whole Pilbara nonulation		20 1.6 390.2 30 2.4 232.2	10
		With reference to the targets, the term 'long-term' needs to be defined (Table 1)		50 4.1 120.8	
				75 6.1 71.9	
				200 16.3 20.5	
				300 24.5 12.2	
				750 61.2 3.8	1
				1000 81.6 2.6	10
			RioTinto	Vibration Prediction for	Peak Parti Scaled Di
				K PPV value when SD = 1 731 mm/s 10 b Local rock exponent -1.28 (-1.6 typ) 10	000
				MIC Max charge in 8ms 150 kg D Distance from blast 350.6 m	
				SD Scaled Distance 28.6 m/vkg	100
				PPV Peak particle velocity 10.0 mm/s	
				10 0.8 947.6	
				20 1.6 390.2	10
				30 2.4 232.2 50 4.1 120.8	
				75 6,1 71.9	
				100 8.2 49.7 200 16.3 20.5	-
				300 24.5 12.2	
				500 40.8 6.3 750 61.2 3.8	1
				1000 81.6 2.6	10
			Figu the s	ıre 2 Initial blast modelling using Brockman Iron geology data fron 95% confidence level	n the Proponent's



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			Blast vibration monitoring will be undertaken for all blasts within 3 100mm/s PPV vibration trigger is not exceeded. This will include monitoring of actual PPV, and analysis of modelled versus actual PPV the Proposal will trigger an update of the mine plan and blast model a as triggered if sooner) inspections for structural damage to the Rat vibration levels exceed 10mm/s PPV.
			Potential impacts to the roost will be managed through reduced vibrat conservative vibration value) and lower call limit (LCL) trigger and thre undertaken continuously with quarterly analysis to identify any char triggered if earlier) inspections of the roost will also be undertaken to
			Infra-red cameras have been used on a periodic basis at the Ratty Sp consistent with most monitoring of PLNB roosts conducted by Rio Tim for calculating the number of calls vs. the number of individual bats to estimated. Infra-red cameras will still periodically be used to readju estimates from the acoustic monitoring.
			Analysis of the long-term data at this roost and other PLNB roosts sug (LCL) is an appropriate monitoring method for identifying potential in the Proposal. Roosts will naturally trigger the LCL, however prolonge decline. The population count for the roost is highly variable both long-term data (monitored since 2015) provided in Figure 8 of Bat Cal based on a LCL that is derived from the long-term average of nightly cal estimated population at the roost is considered too complex as the variation, population movements, relationships to other unknown roo (pers. com Robert Bullen, Bat Call WA).
			An upper call limit (UCL) will show population growth at the Ratty Spri proposed trigger or threshold level requiring management for the Pro- more relevant indicator of potential impacts to PLNB population at Ra number of peak events or a seasonal shift will still be continuously (annually) and provide context to the current status of the Ratty Sprin
			Due to the remote location of the roost and the lack of telemetry technologies real-time analysis is unable to be undertaken. As such, continuous monthe only practical way to effectively monitor the roost and any char monitoring by the Proponent at other Pilbara leaf-nosed bat roost site
			Continuous acoustic monitoring data is available for the Ratty Springs R data), with additional data from 2015 and 2017 available. As minin provides more than the minimum two years of baseline data for this s
			Due to the restricted shape and size of the Ratty Springs roost access Pilbara Leaf-nosed Bat scats is not possible. Discussions with Dr. Biodiversity Conference) who has tried to undertake DNA scat analy methods are unsuitable for this species. Accurate population counts of and infra-red camera counts.

300 m of the Ratty Spring roost to ensure the modelling of PPV prior to blasting, vibration /. Exceedances of 10mm/s PPV associated with and blast management procedures. Biannual (or tty Spring cave entrance will be undertaken if

tion levels (a threshold of 10mm/s PPV, the most eshold values. Lower call limit monitoring will be nges to the roost population. Bi-annual (or as identify structural instability (e.g. rock fall).

prings roost through the life of the monitoring, to. This method is used to calibrate the formula enable the Ratty Springs roost population to be ust the formula to ensure accurate population

ggests the formula for creating a lower call limit npacts to the Ratty Spring roost associated with ed or repeated triggers indicate a more serious between and within years, as shown from the II (2020a). The trigger and threshold criteria are II counts; readjusting the LCL based on the yearly ere are too many unknown variables (seasonal bsts in the area etc.), likewise with a seasonal LCL

ings roost. However, population growth is not a popsal. The Proponent is confident that LCL is a atty Springs Roost. Increases or decreases in the monitored, analysed (quarterly) and reported ag roost.

ology currently available for acoustic recordings, nitoring and quarterly analysis currently provides anges to its population. This is consistent with es.

Roost from 2019 (minimum of four years baseline ag activities are set to commence in 2023, this species.

sing the roost chamber to place mats to collect Linette Umbrello (DBCA-presentation at the rsis for this species has also stated that current can currently be obtained via acoustic recorders

Item	Торіс	DWER Comment	Rio Tinto Response
			General Responses As the management of the Ghost Bat population in Western Range is in the Development Envelope, BACI analysis is not deemed suitable. If if there was a large enough sample size of roosts to act as replicants of the small population size (n=10) and lack of roost site fidelity mean Likewise, there is a lack of other known Pilbara Leaf-nosed Bat roosts in a BACI analysis. Instead monitoring undertaken at several other Ric Pilbara wide comparison of population and seasonal trends to act as r For consistency with other projects the aforementioned terms relation been defined and used in the revised EMP (Rio Tinto 2021a). 'Long term' refers to decades (not years) in duration. Long-term permanent loss of environmental value/s, or where intensive and/or up over decades would be required to restore the environmental value/s term adverse (in terms of impact) have been included in the revised E
Short-	range endemic (SI	RF) invertebrates	
3	Summary	The information provided is insufficient to assess potential impacts to SRE invertebrates. This is largely because identifications of the majority of specimens have not been completed to a level (species/morphospecies or genetic lineages) that allows determination of their distributions, and a comparison of the specimens across the entire development envelope has not been attempted. Consequently, there is insufficient information to determine which, if any, species are restricted to impact zones. Therefore, a high degree of uncertainty remains with respect to the potential for adverse impacts to SREs in the development envelope. Further detail is provided below.	Noted, please refer to responses below. Additional supplementary information is provided in Attachment 2 Endemic invertebrates consolidated impact assessment.
4	Response to submissions – general comment	The comments below reflect the provision of new information in the form of a level 1 survey (Biologic 2021) in the Western Range (WR) area and clarification is required on some points, as discussed below.	Noted, please refer to responses below.
5	RTS No. 16	 The RTS states that there were few sampling points within high and moderate suitability habitats in proposed pit areas, particularly in the Western Range area. An additional survey round (Biologic 2021) has been conducted in the WR area and it has adequately sampled the area. However, the report lacks critical information regarding species identifications and their distributions in relation to impacts areas, that are required to assess and quantify impacts to SREs, and a review of the impacts to SREs using consolidated information from all surveys has not been undertaken. The shortfalls in Biologic (2021) and work required to address them are discussed below. Analysis: The report does not consolidate previous survey data and specimens from previous surveys have not been compared. The majority of potential SRE species (12 out of 20) are not identified to species level so their distributions cannot be determined and therefore potential impacts to SRE species cannot be determined. Additionally, Biologic states that 	 Noted. The Proponent has undertaken further SRE analysis and assessme Attachment 2 and Attachment 3 respectively) including: Consolidation of SRE species and habitat information from Development Envelope, including: a. Alignment of SRE identifications from all previous surfue specimens, taxonomic frameworks, and genetic sequed b. Updated taxonomic and ecological information relevant to potential impact areas; Indicative linear ranges in the local or wider regional and d. Consolidation of SRE habitat mapping and occurrent throughout the Development Envelope; and Occurrence of SRE species in relation to proposed implement environmental Impact Assessment of SRE species values and the Proposal, including direct and indirect impacts, and cumu constraints of available data

based on the continued presence of the species BACI design would be a suitable analysis method or control sites. For Ghost Bats in Western Range ns any values would not be statistically robust. in the vicinity to act as appropriate control sites o Tinto sites for both bat species will allow for a reference sites.

ing to bats (roost, colony and population) have

impact is defined as an impact resulting in a in-proven management intervention, potentially s. The definition for long-term impact and long-IMP (Rio Tinto 2021a).

, Biologic (2021a) Memorandum: Short Range

nent (provided in Biologic 2021a and 2021b, om all surveys completed to date within the rveys (as much as practicable based on available ences); nt to assessment of species distributions relative rea for all SRE species, to the best available data; ence of SRE species in habitat types mapped pact areas. habitat values in relation to potential impacts of lative impacts as much as practicable within the

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		 the Study Area, it is difficult to say how restricted these Potential SRE species will be to the Study Area." A number of species complexes have been found in the study area. for example. 	The Biologic (2021a) report has been updated to incorporate further (Attachment 2).
		 5. A number of species complexes have been found in the study area, for example, the isopod Buddelundia 36 type. A species complex is a taxon that is composed of multiple cryptic (difficult to identify based on morphology) species and usually can only be separated using genetic data. Consequently, their inferred distributions should be interpreted with caution because they do not represent a single species. The presence of numerous species complexes and the largely unresolved specimen identifications creates additional uncertainty around the SRE species distributions and makes assessing impacts to SRE fauna particularly challenging. 4. The report uses a novel approach to assess the likelihood of taxa being restricted to the study area. It classifies the potential SRE taxa collected into three groups: 1) potential SRE likely found beyond study area based on habitat type; 2) potential SREs that appear locally abundant with the potential to be restricted in distribution to the range itself; and 3) Potential SRE invertebrate taxa that were represented by one specimen or collection event only. In this instance, the approach has been incorrectly applied because it focuses on whether the taxa are restricted to the study area and not the impact areas. 	 <u>Analysis</u> Biologic's (2021) report presents data from the Western Austiprevious reports within the wider local area, including Astron This includes species found within 40km of the Proposal. Hidentified past the genus level, the ability to compare specime survey specimens is limited. The Proponent has since undertaken molecular analysis to all of different surveys at the species level. Section 2 and Table data consolidation and alignment following completion of all S Section 2 and Table 2.1 (Biologic 2021a) document the method completion of all SRE work. This includes the approach used to or OTUs described in the memorandum. Sections 2.2.1 - 2.2 groups.
		5. Groups 2 and 3 are considered to be potentially restricted to the study area. The report states that "molecular work would be required to confirm whether taxa are restricted to the study area" (p. 6). Group 3 contains singletons for which "distribution cannot be worked out even with further molecular or taxonomic work unless further specimens are located" (p. 6). However, it may be possible to determine distributions for some species by comparing specimens from earlier surveys. For example, Aname specimens have been collected from three sites over the three different surveys and two of these sites appear to be outside the impact areas. Genetic analysis could allow the specimens to be matched and may determine if they are not restricted to impact areas.	The Proponent has undertaken molecular analysis (DNA barco collected within the Study Area and wider local area from both A total of 96 specimens, 60 from the Biologic (2021b) survey sequenced and this information has allowed for species level units (OTUs) or morpho-species) and assessment of distribution The genetic work has now resolved the majority of identifican Table 2.1 (Biologic 2021a) provides an update of local and reg not practicable to obtain a successful sequence for every in previous studies, but a reasonable attempt has been made to co previous identifications with material collected recently by Bi
		 Habitat assessment: 6. The habitat types are not consistent between the different survey reports or the Environmental Review Document (ERD). For example, Astron has not mapped gorges/gullies in WR, while the Biologic mapping shows numerous gorge/gully habitats that they consider high suitability habitat for SREs. 7. The ERD discusses impacts to high and moderately suitable SRE habitat. However, the Biologic report maps habitat types rather than habitat. A consistent mapping approach is recommended to reduce uncertainty. 	 ability to confirm wider species distributions, although available obtain any possible alignments with sequence material from the sequence of species complexed that exist for several groups and the difficulty of assessing implevel. Biologic 2021a Section 2.1 and Table 2.1 document the Sections 2.2.1 - 2.2.6 provide further detail regarding fauna groups
		 Figures: 8. Figure 4.4 maps the locations of potential SREs but has numerous taxon symbols overlapping so the taxon distributions are not clear and it is difficult to interpret. 9. The figures do not illustrate the areas of impact in relation to the species recorded. 10. A map illustrating the locations of all species from all surveys across the development envelope has not been provided. 	 In the absence of clear delineation within species complexes the Proponent and their consultants have taken a conserrepresenting Potential SRE and have used habitat occupancy a distribution within the Development Envelope. Table 2.2 deta each mapped habitat type. 4. The Proponent notes that the Biologic (2021b) report was no and as such considered fauna and fauna habitat restrictio conceptual disturbance footprint. The fauna grouping approacimpact assessment approach in Biologic 2021a.

work and is attached for DWER's consideration

tralian Museum (WAM) database searches and on (2018) and Bennelongia (2012) (Section 4.1). lowever, as most of these specimens were not ens at the species-level with the Biologic (2021b)

low for further comparison between specimens 2.1 (Biologic 2021a) document the methods of SRE work.

ds of data consolidation and alignment following o resolve as many records as possible to species 2.6 document the approach to individual fauna

oding using the mitochondrial gene COI) of taxa in the Biologic (2021b) and Astron (2018) surveys. y and 36 from Paraburdoo (Astron 2018) were determinations (genetic operational taxonomic ons for the taxa collected at Western Range.

ations at the species (or genetic OTU) level, and gional linear ranges for each taxonomic unit. It is individual, especially given legacy issues from obtain species level OTUs for all groups and align iologic. The lack of genetic sequences from the vas again identified as a data gap that limits the ble regional genetic databases were checked to the wider region.

tes due to the incomplete taxonomic frameworks bacts to these potential SRE fauna at the species he collation of available linear range data, and roups including cryptic species.

(particularly those comprising cryptic species), rvative approach and assumed individuals as as a potential surrogate for understanding their ails the occurrence of each of the species within

ot intended to be an impact assessment report, ons within the Study Area as opposed to the ch in Biologic 2021b has been superseded by the

Item	Торіс	DWER Comment	Rio Tinto Response
			 The fauna grouping approach in Biologic 2021b has been super Biologic 2021a. The Proponent has undertaken molecular and gene COI) to resolve species-level identifications and further Biologic (2021b) and Astron (2018) surveys.
			 Following the consolidation and alignment of SRE fauna record have been recorded from the Development Envelope: Eight species are known to be widespread and are not 20 OTUs (including indeterminate taxa) have uncertair The remaining 32 species/ OTUs are considered Potent current taxonomic/ ecological information.
			Three further taxa (<i>Buddelundia</i> `sp. Biologic-ISOP047`, <i>Crypto</i> `sp. Biologic-CHIL025`) represent new, unique OTUs recorde proposed impact areas. Based on current information, the Development Envelope, although their habitats (Gorge/ Gul mosaic of interconnected rocky/ mountainous habitat that Development Envelope and the wider local area. These Development Envelope and the wider area post development. of the Proposal are unlikely to cause a significant risk to the local area.
			 <u>Habitat Assessment</u> 6. The Astron (2018) and Biologic (2021b) SRE survey areas and a Proponent has updated the Figure 6.4 in the ERD to reflect Attachment 1). Further detail regarding habitat types/assess Figure 2.1 (Biologic 2021a, Attachment 2).
			 Biologic (2021a) Table 3.2 provides an updated assessment of within the Development Envelope across all surveys; high (Gor Low Hills, Riverine, and Drainage), and low value (Alluvial Plai Chapter has been revised to reflect the updated habitat value
			Section 2.3 and Table 2.2 Biologic (2021a) details the occurrenc to the habitat types mapped across the Development Envelop the ERD (Attachment 1) and Figure 2.1 Biologic (2021a) (Attach
			 Noted. Figures have been revised and included in the updated Biologic (2021a) memorandum (Attachment 1 and 2 respect Potential SRE taxon symbols making interpretation clearer.
			 Noted. Figures have been revised and included in the updated Biologic (2021a) memorandum (Figure 2.1) (Attachment 1 and records and the conceptual disturbance footprint.
			10. Noted. Figures have been revised and included in the updated Biologic (2021a) memorandum (Figure 2.1) (Attachment 1 and

erseded by the impact assessment approach in nalysis (DNA barcoding using the mitochondrial investigate species distributions from both the

rds (Biologic 2021a) a total of 60 species/OTUs

t considered SREs.

n distributions.

tial SREs based on the nominal range criteria and

ops 'sp. Biologic-CHIL022', and Geophilomorpha ed only from single sites or a few sites, inside ese taxa have not been detected outside the Ily, Rocky Hills, and Low Hills habitats) form a c is extensive beyond impacts throughout the e habitats will remain connected within the Biologic (2021a) conclude the potential impacts ong-term persistence of these three taxa.

associated habitat mapping do not overlap; the t all SRE survey data (Figure 6.4 is included as sment is included in Section 2.3, Table 2.2 and

f the overall habitat value for SRE taxa recorded rge/Gully), medium (Breakaway/Cliff, Rocky Hill, in, and Stony Plain). The Terrestrial Fauna ERD assessment (Rio Tinto 2021b).

ce of each of the SRE species and OTUs in relation be. These are visually presented on Figure 6.4 in hment 2).

d Terrestrial Fauna ERD Chapter (Figure 6.4) and ctively). Figures now have fewer overlapping

d Terrestrial Fauna ERD Chapter (Figure 6.4) and d 2 respectively). Figures present Potential SRE

d Terrestrial Fauna ERD Chapter (Figure 6.4) and d 2 respectively).

 RTS No. 19 The DWER's submission requested a table that included listing where each taxon was found in results of the first location for each specimen fues DWER'S requests. Appendix 10 of the RKS includes a table with the sample location for each specimen fues undertaken further SRE analysis and assessment (analysis) and assessment of impacts to same the same esticated to impact areas. Table 2.3 in Biologic 2021 presents an up-to-date, concolladd in the Development Invelope; Section 2.0 filliogic 2011 presents an up-to-date, concolladd in the Development Invelope; Section 2.0 filliogic 2012 presents and further investigates and analysis indentifications and protein and information requested. Unreactive distributions and potential impacts to SRE invertee investigates and according to inform the approxemation to the impact areas. as requested. The Proponent has undertaken molecular analysis (block to the investigates propenses to 2012). Therefore, the KT does determined in Atran 0.2013 and Biologi (2021) presents and further investigates propenses to rough and exequired if the signet president and there accurate distributions of species in relations to the impact areas. as recommended in Atran 0.2013 and Biologi (2021). Forefore, it is recommended that genetic analysis because to resolve the issues raised in order to provide tables and execution of the uspecies log provide genetic consultation to the videopter and nearboard that genetic analysis because to recolve the issues raised in order to provide the according the proposal. Work required Do be resolution of the uspect result of the consolution of any kan be and contrate provide and assessment of quarks to soft and the resolution of the uspect result of the resolution of the uspect result of	Item	Торіс	DWER Comment	Rio Tinto Response
 Tourd in relation to the impact sites if Le. inside and/or outside disturbance areas. In Appendix 10 of the XT suchas a table with the sample location for each spectram in a subset set within the Elevenement of the analysis and assessment of impacts to sRE interfloate is limited by the unresolved taxonomy and an alignment of the analysis of the max recent list on the impacts to SRE interfloate is limited by the unresolved taxonomy and an alignment of the analysis (MAA builded). For example, Aname sp. indch. Tab ben collected of morphological or genetic codes where relevant). For example, Aname sp. indch. Tab ben collected of morphological or genetic codes where relevant). For example, Anames p. indch. Tab ben collected of morphological or genetic codes where relevant). For example, Anames p. indch. Tab ben collected of the sciences of th	6	RTS No. 19	The DWER submission requested a table that included listing where each taxon was	The Proponent notes DWER'S requests.
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 2021). Therefore, the RIS does not provide the information required to sesses specifies a number of the information and potential impacts to setting specific as nost of the taxes are not identifications. The assessment of impacts to SRE invertebrates is limited by the unresolved taxonomic identifications in addition to information is additional taxonomic identifications and there invertigates genetics analysis (e.g. genetic analysis (wall by experiments from 2011 survey were that the neutral would likely be too old or not preserved quality sequences (Biologic 2021a). A total of 96 specimens, for form the accurate distributions of species in relation to the impact areas, as recommended in Ataro 2018 and Biologic 2021a). A total of 96 specimens, 60 from the Biologic (2021b) surve sequences (Biologic 2021a). A total of 96 specimens, 60 from the Biologic (2021b) surve sequences and accurate distributions of species in recommended in Ataro 2018 and Biologic 2021a). Work required ff the second that genetic analysis be used to resolve the issues raised in order to provide data of the impact assessment. Resolve the lack of a consolidated report and resolution of the outstanding tax interminate specifics (PA 2015). Therefore, it is recommended that genetic analysis be used to resolve the issues raised in order to provide and path impact assessment. Resolve the lack of a consolidated report and resolution of the outstanding tax described the distributions of species and identify any tax that are potential threats to SREs from the proposal. Biologic (2021a) have since consolidated and a aligned all SR molecular work; this has resulted in a total of 60 species (2021a) identified information. Consolidate for assessment. Revise the habitat segssment and quantify areas of impact to escale and path to and aligned all SR molecular work; this has resulted in a total of 60 species and identifications a			addition, the table does not contain the results of the most recent surveys (Biologic	proposed impact areas.
 Unresolve description electron or concernence on conc			2021). Therefore, the RTS does not provide the information required to assess species distributions and notential impacts to restricted species as requested	Table 2.1 in Biologic 2021a presents an up-to-date, consolidate within the Development Envelope: Section 2 of Biologic
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 and no alignment of the majority of specimens collected from four sites but it is not possible to determine whether it is a single species without further analysis (e.g. gueences analysis or specimens from 2011 survey, server that the material would likely be too old or not preserved quality sequences (Biologic 2021a). Arati of 96 specimens, 60 from the Biologic (2021b) survey sequences (Biologic 2021a). A trai of 96 specimens, 60 from the Biologic (2021b) survey sequences (Biologic 2021a). A trai of 96 specimens, 60 from the Biologic (2021b) survey sequences (Biologic 2021a). A trai of 96 specimens, 60 from the Biologic (2021b) survey sequences (Biologic 2021a). A trai of 96 specimens, 60 from the Biologic (2021b) survey sequences (Biologic 2021a). A trai of 96 specimens, 60 from the Biologic (2021b) survey sequences (Biologic 2021a). A trai of 96 specimens, 60 from the Biologic (2021b) survey sequences (Biologic 2021a). A tread of a consolidated report and resolution of the outstanding tag identifies and their distributions, there is currently insufficient information to assess the potential impact assessment: A resolve the identifications of all taxa collected using genetic analysis, including comparison of specimens, collected from Attron (2018) and Biologic (2021). A resolve the infiftications of all taxa collected using genetic analysis, including comparison of specimens, collected from Attron (2018) and Biologic (2021b) identified limitations and constraints to under a considered Poten current taxonomic level possible or and the information. Consolidate the impact assessment and quantify area so fimpact to acass. Revise the habitat assessment and quantify area so fimpact to acass. Revise the habitat assessment and quantify area so fimpact to acas. Revise the habitat assessment and quantify areas of dearing and the edivinginal and management measures specific for SRE taxa.			The assessment of impacts to SRE invertebrates is limited by the unresolved taxonomy	1. The Proponent has undertaken molecular analysis (DNA ba
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 or will not adequately address questions of SRE restriction to proposed impact areas* (EPA 2016). Therefore, it is recommended that genetic analysis be used to resolve the issues raised in order to provide data for the impact assessment and reduce uncertainty regarding the potential threats to SREs from the proposal. Work required Due to the lack of a consolidated report and resolution of the outstanding tax identities and their distributions, there is currently insufficient information to assess the potential impacts to SREs in the development envelope. The following work is required to enable impact assessment: Resolve the identifications of all taxa collected using genetic analysis, including indeterminate taxa) have uncerta comparison of specimens collected from Astron (2018) and Biologic (2021). Provide an explanation for any taxa that cannot be resolved (e.g. old or poor quality specimen). Consolidate the records from all surveys across the entire development area, describe the distributions of species and identify any taxa that are potentially restricted to the impact areas. Revise the habitat assessment and quantify areas of impact to each SRE habitat type. Provide mapping to illustrate the SRE habitat types in relation to the cumulative impacts to SREs in the development envelope incorporating the results of the consolidated review and discuss the cumulative impacts to SREs from the existing approved areas of clearing and the additional areas as part of this Proposal, as per ESD Item 13b and 16. Discuss mitigation and management measures specific for SRE taxa. 			analysis methods in situations where traditional taxonomic identification is not possible	In total, the melecular analysis resulted in 20 OTUs being d
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 Issues assed in order to provide data tor the impact assessment and reduce uncertainty regarding the potential threats to SREs from the proposal. Work required Due to the lack of a consolidated report and resolution of the outstanding taxa identities and their distributions, there is currently insufficient information to assess the potential impacts to SREs in the development envelope. The following work is required to enable impact assessment: Resolve the identifications of all taxa collected using genetic analysis, including indeterminate taxa) have uncertain comparison of specimens collected from Astron (2018) and Biologic (2021). Provide an explanation for any taxa that cannot be resolved (e.g. old or poor quality specimen). Consolidate the records from all surveys across the entire development area, describe the distributions of species and identify any taxa that are potentially restricted to the impact areas. Revise the habitat assessment and quantify areas of impact to each SRE habitat type. Provide mapping to illustrate the SRE habitat types in relation to the known locations of the taxa (identified to lowest taxonomic level possible) consolidated from all surveys, as per the work required in the Environmental Scoping Document (ESD) (Item 15). Revise the habitat assess the potential impacts to SREs in the development envelope, there will always b cannot be resolved morphologically, genetically, or spatially genetic alusonomic frameworks disagree with genetic a so-sampling mean that not all the material can b contaminated, sometimes the material is too degraded fm orphological taxonomic frameworks disagree with genetic rabe and interet and management measures specific for SRE taxa. 			(EPA 2016). Therefore, it is recommended that genetic analysis be used to resolve the	(2024b) Study Associated and Paraburdoo). From these spec
 Work required Due to the lack of a consolidated report and resolution of the outstanding taxa identities and their distributions, there is currently insufficient information to assess the potential impacts to SREs in the development envelope. The following work is required to enable impact assessment: Resolve the identifications of all taxa collected using genetic analysis, including comparison of specimes collected from Astron (2018) and Biologic (2021). Provide an explanation for any taxa that cannot be resolved (e.g. old or poor quality specimen). Consolidate the records from all surveys across the entire development area, describe the distributions of species and identify any taxa that are potentially restricted to the impact areas. Revise the habitat assessment and quantify areas of impact to each SRE habitat type. Provide mapping to illustrate the SRE habitat types in relation to the known locations of the taxa (identified to lowest taxonomic) are collected in the Development Envelope, there will always be consolidated from all surveys, as per the work required in the Environmental Scoping Document (ESD) (Item 15). Re-analyse and assess the potential impacts to SREs in the development envelope incorporating the results of the consolidated review and discuss the cumulative impacts to SREs from the existing approved areas of clearing and the additional areas as part of this Proposal, as per ESD Item 13b and 16. Discuss mitigation and management measures specific for SRE taxa. 			issues raised in order to provide data for the impact assessment and reduce uncertainty	(2021b) Study Area, not including indeterminate species (sp.
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 Biologic (2021a) have since consolidated and aligned all SR inverting the potential impacts to SREs in the development envelope. The following work is required to enable impact assessment: Resolve the identifications of all taxa collected using genetic analysis, including comparison of specimens collected from Astron (2018) and Biologic (2021). Provide an explanation for any taxa that cannot be resolved (e.g. old or poor quality specimen). Consolidate the records from all surveys across the entire development area, describe the distributions of species and identify any taxa that are potentially restricted to the impact areas of impact to each SRE habitat type. Provide mapping to illustrate the SRE habitat types in relation to the known locations of the taxa (identified to lowest taxonomic level possible consolidated from all surveys, as per the work required in the Environmental Scoping Document (ESD) (Item 15). Re-analyse and assess the potential impacts to SREs in the development envelope incorporating the results of the consolidated review and discuss the cumulative impacts to SREs from the existing approved areas of clearing and the additional areas as part of this Proposal, as per ESD Item 13b and 16. Discuss mitigation and management measures specific for SRE taxa. 			Most required	Confirmed SRE based on data available to date.
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 Interpotential impacts to SREs in the evelopment environmental scoping Document (ESD) (Item 15). Re-analyse and assess the potential impacts to SREs from the existing approved areas of clearing and the additional areas as part of this Proposal, as per ESD Item 13b and 16. Discuss mitigation and management measures specific for SRE taxa. Envelope: Eight species are known to be widespread and are not current taxonomic / ecological information. Envelope: Eight species are known to be widespread and are not current taxonomic / ecological information. Consolidate the records from all surveys across the entire development area, describe the distributions of species and identify any taxa that are potentially restricted to the impact areas. Revise the habitat assessment and quantify areas of impact to each SRE habitat types in relation to the known locations of the taxa (identified to lowest taxonomic level possible) consolidated from all surveys, as per the work required in the Environmental Scoping Document (ESD) (Item 15). Re-analyse and assess the potential impacts to SREs from the existing approved areas of clearing and the additional areas as part of this Proposal, as per ESD Item 13b and 16. Discuss mitigation and management measures specific for SRE taxa. 			the notantial impacts to SPEs in the development envelope. The following work is	molecular work; this has resulted in a total of 60 species
 Resolve the identifications of all taxa collected using genetic analysis, including comparison of specimens collected from Astron (2018) and Biologic (2021). Provide an explanation for any taxa that cannot be resolved (e.g. old or poor quality specimen). Consolidate the records from all surveys across the entire development area, describe the distributions of species and identify any taxa that are potentially restricted to the impact areas. Revise the habitat assessment and quantify areas of impact to each SRE habitat type. Provide mapping to illustrate the SRE habitat types in relation to the known locations of the taxa (identified to lowest taxonomic level possible) consolidated from all surveys, as per the work required in the Environmental Scoping Document (ESD) (Item 15). Re-analyse and assess the potential impacts to SREs in the development envelope incorporating the results of the consolidated review and discuss the cumulative impacts to SREs from the existing approved areas of clearing and the additional areas as part of this Proposal, as per ESD Item 13b and 16. Discuss mitigation and management measures specific for SRE taxa. The Proponent has undertaken further work which is present the Development further work which is present the Development further work disagree with genetic resoluted of all SRE invert the Development further work disagree with genetic resoluted and the additional areas as part of this Proposal, as per ESD Item 13b and 16. The Proponent has undertaken further work which is present of Table 2.1 provides a consolidated list of all SRE invert the Development further work which is present the Development fu			required to enable impact assessment:	Envelope:
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 The remaining 32 species/ OTUs are considered Poten current taxonomic/ ecological information. Consolidate the records from all surveys across the entire development area, describe the distributions of species and identify any taxa that are potentially restricted to the impact areas. Revise the habitat assessment and quantify areas of impact to each SRE habitat type. Provide mapping to illustrate the SRE habitat types in relation to the known locations of the taxa (identified to lowest taxonomic level possible) consolidated from all surveys, as per the work required in the Environmental Scoping Document (ESD) (tem 15). Re-analyse and assess the potential impacts to SREs in the development envelope incorporating the results of the consolidated review and discuss the cumulative impacts to SREs from the existing approved areas of clearing and the additional areas as part of this Proposal, as per ESD Item 13b and 16. Discuss mitigation and management measures specific for SRE taxa. The Proponent has undertaken further work which is present the Development Envelope 			comparison of specimens collected from Astron (2018) and Biologic (2021)	 20 OTUs (including indeterminate taxa) have uncertain
 current taxonomic/ ecological information. Biologic (2021b) identified limitations and constraints to under a variable to undertake comparisons The accessibility of pre-existing regional sequences The success rate of genetic sequencing. Biologic (2021a) note that despite the successful taxonomic al collected in the Development Envelope, there will always be cannot be resolved morphologically, genetically, or spatially genetic sub-sampling mean that not all the material is too degraded f morphological taxonomic frameworks disagree with genetic resoluted is present the 2. The Proponent has undertaken further work which is present the Development Envelope			Provide an explanation for any taxa that cannot be resolved (e.g. old or noor	The remaining 32 species/ OTUs are considered Poten
 Consolidate the records from all surveys across the entire development area, describe the distributions of species and identify any taxa that are potentially restricted to the impact areas. Revise the habitat assessment and quantify areas of impact to each SRE habitat type. Provide mapping to illustrate the SRE habitat types in relation to the known locations of the taxa (identified to lowest taxonomic level possible) consolidated from all surveys, as per the work required in the Environmental Scoping Document (ESD) (Item 15). Re-analyse and assess the potential impacts to SREs in the development envelope incorporating the results of the consolidated review and discuss the cumulative impacts to SREs from the existing approved areas of clearing and the additional areas as part of this Proposal, as per ESD Item 13b and 16. Discuss mitigation and management measures specific for SRE taxa. 			quality specimen)	current taxonomic/ ecological information.
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 3. Revise the habitat assessment and quantify areas of impact to each SRE habitat type. Provide mapping to illustrate the SRE habitat types in relation to the known locations of the taxa (identified to lowest taxonomic level possible) consolidated from all surveys, as per the work required in the Environmental Scoping Document (ESD) (Item 15). 4. Re-analyse and assess the potential impacts to SREs in the development envelope incorporating the results of the consolidated review and discuss the cumulative impacts to SREs from the existing approved areas of clearing and the additional areas as part of this Proposal, as per ESD Item 13b and 16. 5. Discuss mitigation and management measures specific for SRE taxa. 4. The Proponent has undertaken further work which is present - Table 2.1 provides a consolidated list of all SRE invert the Development Envelope 			restricted to the impact areas.	 Breadth of data available to undertake comparisons
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 known locations of the taxa (identified to lowest taxonomic level possible) consolidated from all surveys, as per the work required in the Environmental Scoping Document (ESD) (Item 15). 4. Re-analyse and assess the potential impacts to SREs in the development envelope incorporating the results of the consolidated review and discuss the cumulative impacts to SREs from the existing approved areas of clearing and the additional areas as part of this Proposal, as per ESD Item 13b and 16. 5. Discuss mitigation and management measures specific for SRE taxa. 8. Biologic (2021a) note that despite the successful taxonomic and collected in the Development Envelope, there will always be cannot be resolved morphologically, genetically, or spatially or genetic sub-sampling mean that not all the material can be contaminated, sometimes the material is too degraded from orphological taxonomic frameworks disagree with genetic resolved morphological taxonom			type. Provide mapping to illustrate the SRE habitat types in relation to the	 The success rate of genetic sequencing.
 consolidated from all surveys, as per the work required in the Environmental Scoping Document (ESD) (Item 15). Re-analyse and assess the potential impacts to SREs in the development envelope incorporating the results of the consolidated review and discuss the cumulative impacts to SREs from the existing approved areas of clearing and the additional areas as part of this Proposal, as per ESD Item 13b and 16. Discuss mitigation and management measures specific for SRE taxa. Biologic (2021a) note that despite the successful taxonomic and collected in the Development Envelope, there will always be cannot be resolved morphologically, genetically, or spatially of genetic sub-sampling mean that not all the material can be contaminated, sometimes the material is too degraded from orphological taxonomic frameworks disagree with genetic resolved morphological taxonomic			known locations of the taxa (identified to lowest taxonomic level possible)	
Scoping Document (ESD) (Item 15). 4. Re-analyse and assess the potential impacts to SREs in the development envelope incorporating the results of the consolidated review and discuss the cumulative impacts to SREs from the existing approved areas of clearing and the additional areas as part of this Proposal, as per ESD Item 13b and 16. 5. Discuss mitigation and management measures specific for SRE taxa. 2. The Proponent has undertaken further work which is present • Table 2.1 provides a consolidated list of all SRE invert the Development Envelope			consolidated from all surveys, as per the work required in the Environmental	Biologic (2021a) note that despite the successful taxonomic ar
 4. Re-analyse and assess the potential impacts to SREs in the development envelope incorporating the results of the consolidated review and discuss the cumulative impacts to SREs from the existing approved areas of clearing and the additional areas as part of this Proposal, as per ESD Item 13b and 16. 5. Discuss mitigation and management measures specific for SRE taxa. 2. The Proponent has undertaken further work which is present the Development Envelope 			Scoping Document (ESD) (Item 15).	collected in the Development Envelope, there will always be
 envelope incorporating the results of the consolidated review and discuss the cumulative impacts to SREs from the existing approved areas of clearing and the additional areas as part of this Proposal, as per ESD Item 13b and 16. 5. Discuss mitigation and management measures specific for SRE taxa. 2. The Proponent has undertaken further work which is present Table 2.1 provides a consolidated list of all SRE invert the Development Envelope 			4. Re-analyse and assess the potential impacts to SREs in the development	cannot be resolved morphologically, genetically, or spatially
cumulative impacts to SREs from the existing approved areas of clearing and the additional areas as part of this Proposal, as per ESD Item 13b and 16. 5. Discuss mitigation and management measures specific for SRE taxa. 5. The Proponent has undertaken further work which is present • Table 2.1 provides a consolidated list of all SRE invert the Development Envelope			envelope incorporating the results of the consolidated review and discuss the	genetic sub-sampling mean that not all the material can be
the additional areas as part of this Proposal, as per ESD Item 13b and 16. 5. Discuss mitigation and management measures specific for SRE taxa. 2. The Proponent has undertaken further work which is present • Table 2.1 provides a consolidated list of all SRE invert the Development Envelope			cumulative impacts to SREs from the existing approved areas of clearing and	contaminated cometimes the material is too degraded f
 5. Discuss mitigation and management measures specific for SRE taxa. 2. The Proponent has undertaken further work which is present Table 2.1 provides a consolidated list of all SRE invert the Development Envelope 			the additional areas as part of this Proposal, as per ESD Item 13b and 16.	morphological taxonomic frameworks disagree with genetics
 2. The Proponent has undertaken further work which is present Table 2.1 provides a consolidated list of all SRE invert the Development Envelope 			5. Discuss mitigation and management measures specific for SRE taxa.	
Table 2.1 provides a consolidated list of all SRE invert the Development Envelope				2. The Proponent has undertaken further work which is present
the Development Envelope				 Table 2.1 provides a consolidated list of all SRE inverted
				the Development Envelope

Biologic 2021a and 2021b) including: ated list of taxa and occurrence in relation to

ed list of species identifications across all surveys 2021a provides detail on the alignment and on provided in Biologic 2021b.

arcoding using the mitochondrial gene COI) to ecies distributions from both the Biologic (2021b) not chosen for genetic analysis as it was deemed adequately and hence unlikely to produce high

y and 36 from Paraburdoo (Astron 2018) were ies level determinations (genetic OTUs/morphouence data exists.

esignated to specimens from the Development imens, 23 taxa were identified from the Biologic indet.), of which 16 were regarded as Potential 'sp. Biologic-ARAN036` was considered to be a

E fauna records including consideration of the /OTUs being recorded from the Development

t considered SREs.

- n distributions.
- tial SREs based on the nominal range criteria and

ertaking molecular analysis to include:

nd genetic alignment of most of the SRE material e a residual amount of specimen material that due to a variety of reasons. The practicalities of e sequenced, some DNA sequences fail or are for genetic or morphological work, and often results (Biologic 2021a).

ed in Biologic (2021a) (Attachment 2), including: ebrate fauna recorded from all surveys within

Item	Торіс	DWER Comment	Rio Tinto Response
			• Table 2.2 details the occurrence of each of the SRE
			mapped across the Development Envelope
			 Table 2.3 presents taxa distribution in relation to th
			• This data is visually presented in Figure 2.1.
			Following the consolidation and alignment of SRE fauna re
			have been recorded from the Development Envelope as de
			Three taxa (Buddelundia `sp. Biologic-ISOP047`, Cryptops
			Biologic-CHIL025`) represent new, unique OTUs recorded or
			impact areas. Based on current information, these taxa h
			Envelope, although their habitats (Gorge/ Gully, Rocky
			interconnected rocky/ mountainous habitat that is extensi
			Envelope and the wider local area. Biologic (2021a) conclud
			to cause a significant risk to the long-term persistence of th
			3. The Proponent has revised the habitat assessment including
			2):
			 Section 5.2 - 5.5 presents potential direct impacts to Table 2.2 datails the occurrence of each of the SPE
			 Table 2.2 details the occurrence of each of the SKE manned across the Development Envelope
			Table 3.2 presents habitat type and proportion (
			following implementation of the Proposal
			• This data is visually presented on Figure 2.1.
			The Terrestrial Fauna ERD Chapter has also been updated alo
			habitats and SRE taxa have been shown based on all relev
			resolved to species or genetic OTU level). None of the habi
			Development Envelope, and all habitats are known to occur
			 The Proponent has considered potential direct and indirect of all survey data.
			Most species and OTUs (53 out of 60 taxa) were recorded b
			outside of impacts (Biologic 2021a). The SRE habitat type
			extensively beyond proposed impact footprint, with a high p
			proposed impact areas. The potential impacts to these 53 to
			the Proposal were therefore considered Low.
			Four indeterminate taxa were recorded from sites with fro
			and proposed infrastructure areas. In each of these four ca
			habitat was minimal, and suitable habitat is likely to remain
			Biologic (2021a) assessed the potential impacts to these fou
			be Low.
			Three further taxa (<i>Buddelundia</i> `sp. Biologic-ISOF
			Geophilomorpha sp. Biologic-CHIL025') represent new, uni
			sites, inside proposed impact areas (Biologic 2021a). Based
1	1		detected outside the Development Envelope, although the

ecies and OTUs in relation to the habitat types

conceptual disturbance footprint impacts

rds (Biologic 2021a) a total of 60 species/OTUs ibed above under item 1 of this RTS No. 19.

p. Biologic-CHIL022`, and Geophilomorpha `sp. from single sites or a few sites, inside proposed e not been detected outside the Development lls, and Low Hills habitats) form a mosaic of e beyond impacts throughout the Development he potential impacts of the Proposal are unlikely e three taxa.

letail presented in Biologic (2021a) (Attachment

REs and SRE habitat pecies and OTUs in relation to the habitat types

total area (%) likely to be disturbed/retained

g with associated figures. Relationships between t sampling records (those that were able to be t types are expected to be strictly limited to the nore widely in the local and regional area.

mulative impacts to SREs following consolidation

yond proposed impact areas, or both inside and that these taxa are associated with also occur portion of their current extent occurring outside a and their habitats from the implementation of

existing infrastructure impacts or from existing s, the combined disturbance to the surrounding ntact, supporting the persistence of SRE species. axa from the implementation of the Proposal to

47`, *Cryptops* `sp. Biologic-CHILO22`, and ne OTUs recorded only from single sites or a few n current information, these taxa have not been habitats are well represented in the wider local

7 BSA Please note that any survey reports or data that are revised after their initial area beyond the Development Envelope. Interconnected or data where application of the revised ERD Terres 7 BSA Please note that any survey reports or data that are revised after their initial Noted.	Item	Торіс	DWER Comment	Rio Tinto Response
7 IBSA Please note that any survey reports or data that are revised after their initial acceptance into the Index of Biodiversity Surveys for Assessments (IBSA). The Proponent should create the sessment process – should be updated in IBSA. The Proponent should create the sessment for the sessest and data where approximation of the filling and construction areas 7 IBSA Please note that any survey reports or data that are revised after their initial acceptance into the Index of Biodiversity Surveys for Assessments (IBSA). The Proponent should create the sessement (IBSA) to facilitate assessment to fees the addition and the sessement (IBSA). The proponent should create the	Item	Topic	DWER Comment	Rio Tinto Responsearea beyond the Development Envelope. The SRE habitats (Gform a mosaic of interconnected rocky/ mountainous habitat toDevelopment Envelope and the wider local area. The residuehabitat from development of the Proposal are Low. BiologicProposal to be unlikely to cause a significant risk to the lowDevelopment Envelope.Although the Proposal may result in the direct loss of SRE incomposed changes at sites within the impact footprint, these impact footprint, these impact footprint is because the recorded occomposed to the sampling artefacts inherent in SRE surveys. Especially for single sites, it is unlikely that a single sampling location would species; rather, it indicates an association to a particular hab the sampling to date failed to detect a more complete distributes of the sampling to date failed to detect a more complete distributes of the sampling to date failed to detect a more complete distributes of the sampling to date failed to detect a more complete distributes of the sampling to date failed to detect a more complete distributes of the sampling to date failed to detect a more complete distributes of the sampling to date failed to detect a more complete distributes of the sampling to date failed to detect a more complete distributes of the sampling to date failed to detect a more complete distributes of the sampling to date failed to detect a more complete distributes of the sampling to date failed to detect a more complete distributes of the sampling to date failed to detect a more complete distributes of the sampling to date failed to detect a more complete distributes of the sampling to date failed to detect a more complete distributes of the sampling to date failed to detect a more complete distributes of the sampling to date failed to detect a more complete distributes of the sampling to date failed to detect a more complete distributes of
7 IBSA Please note that any survey reports or data that are revised after their initial acceptance into the Index of Biodiversity Surveys for Assessments (IBSA) – e.g. as a result of the assessment process – should be updated in IBSA. The Proponent should the Proponent will provide revised survey reports and data where app contact ibsa@dwer wa gov au for assistance in such cases Noted.				 Proposal. 5. The proposed impact areas have been revised to avoid impacts Gorge/Gully and Riverine habitat as much as practicable fo (avoidance, minimisation, rehabilitation, offset). Mining Excluse Rocky Hills, and Low Hills habitats will help mitigate impacts to measures are outlined in Section 6.7 of the revised ERD Terrer a focus on high value fauna habitats. It is considered that additional mitigation and management methe ERD are not necessary due to: the relatively limited spatial extent of the proposed in the known or likely wider occurrence of all SRE habitat Development Envelope the minor effects of indirect impacts that are likely liming and construction areas the application of the mitigation hierarchy to manage detailed in section 6.7 of the Terrestrial Fauna ERD Chemister
acceptance into the Index of Biodiversity Surveys for Assessments (IBSA) – e.g. as a result of the assessment process – should be updated in IBSA. The Proponent should contact ibsa@dwar wa gov au for assistance in such cases	7	IBSA	Please note that any survey reports or data that are revised after their initial	Noted.
contact ibsa@dwar wa gov au for assistance in such cases			acceptance into the Index of Biodiversity Surveys for Assessments (IBSA) – e.g. as a	The Propenent will provide revised survey reports and data where any
			contact ibsa@dwer.wa.gov.au for assistance in such cases.	Assessments (IBSA) to facilitate assessment.

Gorge/ Gully, Rocky Hills, and Low Hills habitats) that is extensive beyond impacts throughout the al impacts to Rocky Hills habitat and Low Hills c (2021a) assessed the potential impacts of the ng-term persistence of these three taxa in the

dividuals via mining activities, clearing, and land bacts are unlikely to cause the loss of any species currence of SRE species and OTUs within the e or distribution of any of the taxa recorded, due or the singletons and species recorded only from d encapsulate the total distribution range of the itat type that can be assessed as a proxy where ution range.

sment of potential impacts associated with the

s to high value SRE habitats including Breakaway, Ilowing application of the mitigation hierarchy sions Zones (MEZ) with coverage of Gorge/ Gully, high value habitat. Mitigation and management estrial Fauna ERD Chapter (Rio Tinto 2021b) with

neasures (Biologic 2021a) to those proposed in

mpacts to SRE habitat at types in the local area beyond the

nited to the immediate vicinity of the proposed

known residual impacts as far as practicable as napter.

propriate for the Index of Biodiversity Surveys for

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Rio Tinto. 2021a. Environmental Management Plan, Greater Paraburdoo Iron Ore Hub. December 2021.

Rio Tinto. 2021b. Environmental Review Document – Terrestrial Fauna. Rio Tinto.

Attachment 1

Figure 6.4, Environmental Review Document – Terrestrial Fauna. Rio Tinto. 2021b.

12









Attachment 2

Biologic Environmental Survey. 2021a. *Memorandum: Greater Paraburdoo Short-Range Endemic Invertebrate Fauna Consolidated Impact Assessment*. Report to Rio Tinto Iron Ore. December 2021.

Greater Paraburdoo Iron Ore Hub Proposal (Assessment No. 2189)

17

Attachment 3

Biologic Environmental Survey. 2021b. Western Range Project Short-Range Endemic Invertebrate Fauna Survey. Report to Rio Tinto Iron Ore. December 2021.

18