Table S1. Sampling information for strains genotyped for microsatellite markers, and sequenced at the mitochondrial loci *ND4* and *ND6L.*

|  |  |  |  |  |  |  |
| --- | --- | --- | --- | --- | --- | --- |
| Strain ID | Sampling locality | GPS | Host beetle species | Beetle ID | Msats genotyped? | MtDNA fragment sequenced? |
| RS5340 | BV | -21.34N, 55.71E | *Adoretus* sp. | AdoBV\_3 |  | Y |
| RS5341 | BV | -21.34N, 55.71E | *Adoretus* sp. | AdoBV\_1 |  | Y |
| RS5342 | BV | -21.34N, 55.71E | *Adoretus* sp. | AdoBV\_2 | Y | Y |
| RS5406 | BV | -21.34N, 55.71E | *Adoretus* sp. | AdoBV\_1 | Y | Y |
| RSA037 | BV | -21.34N, 55.71E | *Adoretus* sp. | AdoBV\_6 | Y | Y |
| RSA038 | BV | -21.34N, 55.71E | *Adoretus* sp. | AdoBV\_7 | Y | Y |
| RSA039 | BV | -21.34N, 55.71E | *Adoretus* sp. | AdoBV\_11 | Y | Y |
| RSA040 | BV | -21.34N, 55.71E | *Adoretus* sp. | AdoBV\_18 | Y | Y |
| RSA041 | BV | -21.34N, 55.71E | *Adoretus* sp. | AdoBV\_23 |  | Y |
| RSA042 | BV | -21.34N, 55.71E | *Adoretus* sp. | AdoBV\_35 | Y | Y |
| RSA043 | BV | -21.34N, 55.71E | *Adoretus* sp. | AdoBV\_41 |  | Y |
| RSA044 | BV | -21.34N, 55.71E | *Adoretus* sp. | AdoBV\_41 | Y | Y |
| RSA045 | BV | -21.34N, 55.71E | *Adoretus* sp. | AdoBV\_41 | Y | Y |
| RSB001 | CC | -21.21N, 55.64E | *Amneidus godefroyi* | AmCC\_1 | Y | Y |
| RSB002 | CC | -21.21N, 55.64E | *Amneidus godefroyi* | AmCC\_1 | Y | Y |
| RSB003 | CC | -21.21N, 55.64E | *Amneidus godefroyi* | AmCC\_1 | Y | Y |
| RSB004 | CC | -21.21N, 55.64E | *Amneidus godefroyi* | AmCC\_1 | Y | Y |
| RSB005 | CC | -21.21N, 55.64E | *Amneidus godefroyi* | AmCC\_6 | Y | Y |
| RSB006 | CC | -21.21N, 55.64E | *Amneidus godefroyi* | AmCC\_6 | Y | Y |
| RSB007 | CC | -21.21N, 55.64E | *Amneidus godefroyi* | AmCC\_6 | Y | Y |
| RSB008 | CC | -21.21N, 55.64E | *Amneidus godefroyi* | AmCC\_7 | Y | Y |
| RSB009 | CC | -21.21N, 55.64E | *Amneidus godefroyi* | AmCC\_7 | Y | Y |
| RSB010 | CC | -21.21N, 55.64E | *Amneidus godefroyi* | AmCC\_7 | Y | Y |
| RSB011 | CC | -21.21N, 55.64E | *Amneidus godefroyi* | AmCC\_7 | Y | Y |
| RSB012 | CC | -21.21N, 55.64E | *Amneidus godefroyi* | AmCC\_7 | Y | Y |
| RSB013 | CC | -21.21N, 55.64E | *Amneidus godefroyi* | AmCC\_11 | Y | Y |
| RSB014 | CC | -21.21N, 55.64E | *Amneidus godefroyi* | AmCC\_11 | Y | Y |
| RSB015 | CC | -21.21N, 55.64E | *Amneidus godefroyi* | AmCC\_11 | Y | Y |
| RSB016 | CC | -21.21N, 55.64E | *Amneidus godefroyi* | AmCC\_11 | Y | Y |
| RSB017 | CO | -20.91N, 55.42E | *Adoretus* sp. | AdoCO\_1 | Y | Y |
| RSB018 | CO | -20.91N, 55.42E | *Adoretus* sp. | AdoCO\_1 | Y | Y |
| RSB019 | CO | -20.91N, 55.42E | *Adoretus* sp. | AdoCO\_1 | Y | Y |
| RSB020 | CO | -20.91N, 55.42E | *Adoretus* sp. | AdoCO\_1 | Y | Y |
| RSB021 | CO | -20.91N, 55.42E | *Adoretus* sp. | AdoCO\_1 | Y | Y |
| RSB023 | CO | -20.91N, 55.42E | *Hoplochelus sp.* | HchCO\_3 | Y | Y |
| RSB024 | CO | -20.91N, 55.42E | *Hoplochelus sp.* | HchCO\_3 | Y | Y |
| RSB025 | CO | -20.91N, 55.42E | *Hoplochelus sp.* | HchCO\_3 | Y | Y |
| RSB026 | CO | -20.91N, 55.42E | *Hoplochelus sp.* | HchCO\_3 | Y | Y |
| RSB027 | CO | -20.91N, 55.42E | *Hoplochelus sp.* | HchCO\_5 | Y | Y |
| RSB028 | CO | -20.91N, 55.42E | *Hoplochelus sp.* | HchCO\_5 | Y | Y |
| RSB029 | CO | -20.91N, 55.42E | *Hoplochelus sp.* | HchCO\_5 | Y | Y |
| RS5344 | ES | -21.28N, 55.35E | *Maladera affinis* | MalES\_83 | Y | Y |
| RS5345 | ES | -21.28N, 55.35E | *Maladera affinis* | MalES\_85 |  | Y |
| RS5350 | ES | -21.28N, 55.35E | *Hoplia retusa* | HopES\_2 | Y | Y |
| RSB030 | FPI | -21.31N, 55.60E | *Adoretus* sp. | AdoFPI\_10 | Y | Y |
| RSB031 | FPI | -21.31N, 55.60E | *Adoretus* sp. | AdoFPI\_18 | Y | Y |
| RSB032 | FPI | -21.31N, 55.60E | *Adoretus* sp. | AdoFPI\_18 | Y | Y |
| RSB067 | FPI | -21.31N, 55.60E | *Adoretus* sp. | AdoFPI\_18 | Y | Y |
| RS5407 | GE | -21.10N, 55.65E | *Adoretus* sp. | AdoGE\_1 | Y | Y |
| RS5408 | GE | -21.10N, 55.65E | *Adoretus* sp. | AdoGE\_2 | Y | Y |
| RS5409 | GE | -21.10N, 55.65E | *Adoretus* sp. | AdoGE\_3 | Y | Y |
| RS5410 | GE | -21.10N, 55.65E | *Adoretus* sp. | AdoGE\_4 | Y | Y |
| RS5411 | GE | -21.10N, 55.65E | *Adoretus* sp. | AdoGE\_5 | Y | Y |
| RSA046 | GE | -21.10N, 55.65E | *Adoretus* sp. | AdoGE\_6 | Y | Y |
| RSA047 | GE | -21.10N, 55.65E | *Adoretus* sp. | AdoGE\_7 | Y | Y |
| RSA048 | GE | -21.10N, 55.65E | *Adoretus* sp. | AdoGE\_7 | Y | Y |
| RSA049 | GE | -21.10N, 55.65E | *Adoretus* sp. | AdoGE\_9 |  | Y |
| RSA050 | GE | -21.10N, 55.65E | *Adoretus* sp. | AdoGE\_10 | Y | Y |
| RSA051 | GE | -21.10N, 55.65E | *Adoretus* sp. | AdoGE\_10 | Y | Y |
| RSA052 | GE | -21.10N, 55.65E | *Adoretus* sp. | AdoGE\_14 |  | Y |
| RSA053 | GE | -21.10N, 55.65E | *Adoretus* sp. | AdoGE\_16 |  | Y |
| RSA054 | GE | -21.10N, 55.65E | *Adoretus* sp. | AdoGE\_17 |  | Y |
| RSA055 | GE | -21.10N, 55.65E | *Adoretus* sp. | AdoGE\_17 | Y | Y |
| RSA056 | GE | -21.10N, 55.65E | *Adoretus* sp. | AdoGE\_17 | Y | Y |
| RSA057 | GE | -21.10N, 55.65E | *Adoretus* sp. | AdoGE\_18 | Y | Y |
| RSA058 | GE | -21.10N, 55.65E | *Adoretus* sp. | AdoGE\_18 |  | Y |
| RSA059 | GE | -21.10N, 55.65E | *Adoretus* sp. | AdoGE\_18 | Y | Y |
| RSA118 | GE | -21.10N, 55.65E | *Chrysomelidae* sp. | ChryGE\_1 | Y | Y |
| RS5431 | LS | -21.04N, 55.15E | Soil | NA | Y | Y |
| RS5361 | NB | -21.19N, 55.64E | Soil | NA | Y | Y |
| RSA075 | NB | -21.19N, 55.64E | *Amneidus godefroyi* | AmNB\_2 | Y | Y |
| RSA076 | NB | -21.19N, 55.64E | *Amneidus godefroyi* | AmNB\_2 | Y | Y |
| RSB033 | NB | -21.19N, 55.64E | *Amneidus godefroyi* | AmNB\_4 | Y | Y |
| RSB034 | NB | -21.19N, 55.64E | *Amneidus godefroyi* | AmNB\_4 | Y | Y |
| RSB035 | NB | -21.19N, 55.64E | *Amneidus godefroyi* | AmNB\_4 | Y | Y |
| RSB036 | NB | -21.19N, 55.64E | *Amneidus godefroyi* | AmNB\_4 | Y | Y |
| RSB037 | NB | -21.19N, 55.64E | *Amneidus godefroyi* | AmNB\_4 | Y | Y |
| RSB038 | NB | -21.19N, 55.64E | *Marronus borbonicus* | MarNB\_6 | Y | Y |
| RSB039 | NB | -21.19N, 55.64E | *Marronus borbonicus* | MarNB\_6 | Y | Y |
| RSB040 | NB | -21.19N, 55.64E | *Marronus borbonicus* | MarNB\_6 | Y | Y |
| RSB041 | NB | -21.19N, 55.64E | *Marronus borbonicus* | MarNB\_6 | Y | Y |
| RSB042 | NB | -21.19N, 55.64E | *Marronus borbonicus* | MarNB\_6 | Y | Y |
| RSA077 | PC | -21.18N, 55.58E | *Hoplia retusa* | HopPC\_2 | Y | Y |
| RSA078 | PC | -21.18N, 55.58E | *Hoplia retusa* | HopPC\_9 | Y | Y |
| RSA079 | PC | -21.18N, 55.58E | *Hoplia retusa* | HopPC\_16 | Y | Y |
| RSA080 | PC | -21.18N, 55.58E | *Hoplia retusa* | HopPC\_26 | Y | Y |
| RSA081 | PC | -21.18N, 55.58E | *Hoplia retusa* | HopPC\_28 | Y | Y |
| RSA082 | PC | -21.18N, 55.58E | *Hoplia retusa* | HopPC\_31 | Y | Y |
| RSA083 | PC | -21.18N, 55.58E | *Hoplia retusa* | HopPC\_33 | Y | Y |
| RSA085 | PC | -21.17N, 55.56E | *Hoplia retusa* | HopPCD\_4 | Y | Y |
| RSA086 | PC | -21.17N, 55.56E | *Hoplia retusa* | HopPCD\_6 | Y | Y |
| RSA087 | PC | -21.17N, 55.56E | *Hoplia retusa* | HopPCD\_12 | Y | Y |
| RSA088 | PC | -21.17N, 55.56E | *Hoplia retusa* | HopPCD\_16 |  | Y |
| RSA120 | PC | -21.18N, 55.58E | *Hoplia retusa* | HopPC\_4 | Y | Y |
| RSB044 | PC | -21.18N, 55.58E | *Hoplia retusa* | HopPC\_29 | Y | Y |
| RSB045 | PC | -21.18N, 55.58E | *Hoplia retusa* | HopPC\_29 | Y | Y |
| RSB046 | PC | -21.18N, 55.58E | *Hoplia retusa* | HopPC\_29 | Y | Y |
| RSB047 | PC | -21.18N, 55.58E | *Hoplia retusa* | HopPC\_29 | Y | Y |
| RSB048 | PC | -21.18N, 55.58E | *Hoplia retusa* | HopPC\_29 | Y | Y |
| RSB049 | PC | -21.18N, 55.58E | *Hoplia retusa* | HopPC\_37 | Y | Y |
| RSB050 | PC | -21.18N, 55.58E | *Hoplia retusa* | HopPC\_37 | Y | Y |
| RSB051 | PC | -21.18N, 55.58E | *Hoplia retusa* | HopPC\_37 | Y | Y |
| RSB052 | PC | -21.18N, 55.58E | *Hoplia retusa* | HopPC\_37 | Y | Y |
| RSB053 | PC | -21.18N, 55.58E | *Hoplia retusa* | HopPC\_37 | Y | Y |
| RSB054 | PC | -21.18N, 55.58E | *Hoplia retusa* | HopPC\_104 | Y | Y |
| RSB055 | PC | -21.18N, 55.58E | *Hoplia retusa* | HopPC\_104 | Y |  |
| RSB056 | PC | -21.18N, 55.58E | *Hoplia retusa* | HopPC\_104 | Y | Y |
| RSB057 | PC | -21.18N, 55.58E | *Hoplia retusa* | HopPC\_104 | Y | Y |
| RSB058 | PL | -21.03N, 55.62E | *Adoretus* sp. | AdoPL\_1 | Y |  |
| RSB059 | PL | -21.03N, 55.62E | *Adoretus* sp. | AdoPL\_1 | Y | Y |
| RSB060 | PL | -21.03N, 55.62E | *Adoretus* sp. | AdoPL\_1 | Y | Y |
| RSB061 | PL | -21.03N, 55.62E | *Adoretus* sp. | AdoPL\_1 | Y | Y |
| RSB062 | PL | -21.03N, 55.62E | *Adoretus* sp. | AdoPL\_3 | Y | Y |
| RSB063 | PL | -21.03N, 55.62E | *Adoretus* sp. | AdoPL\_3 | Y | Y |
| RSB064 | PL | -21.03N, 55.62E | *Adoretus* sp. | AdoPL\_3 | Y | Y |
| RSB065 | PL | -21.03N, 55.62E | *Adoretus* sp. | AdoPL\_3 | Y | Y |
| RSB066 | PL | -21.03N, 55.62E | *Adoretus* sp. | AdoPL\_3 | Y |  |
| RSB076 | PL | -21.03N, 55.62E | *Adoretus* sp. | AphSB\_2 | Y | Y |
| RSA061 | PR | -21.13N, 55.27E | *Hoplochelus sp.* | HchPR\_2 |  | Y |
| RSA094 | RFT | -21.08N, 55.36E | *Oryctes borbonicus* | OryRFT\_1 | Y | Y |
| RSA095 | RFT | -21.08N, 55.36E | *Oryctes borbonicus* | OryRFT\_3 | Y | Y |
| RS5343 | RG | -20.89N, 55.51E | *Alissonotum* sp. | AliRG\_73 |  | Y |
| RS5412 | SB | -21.06N, 55.73E | *Adoretus* sp. | AdoSB\_4 |  | Y |
| RS5413 | SB | -21.06N, 55.73E | *Adoretus* sp. | AdoSB\_14 | Y | Y |
| RS5414 | SB | -21.06N, 55.73E | *Maladera affinis* | MalSB\_1 | Y | Y |
| RS5415 | SB | -21.06N, 55.73E | *Maladera affinis* | MalSB\_2 | Y | Y |
| RS5416 | SB | -21.06N, 55.73E | *Maladera affinis* | MalSB\_4 | Y | Y |
| RS5417 | SB | -21.06N, 55.73E | *Maladera affinis* | MalSB\_10 | Y | Y |
| RS5418 | SB | -21.06N, 55.73E | *Maladera affinis* | MalSB\_23 | Y | Y |
| RS5419 | SB | -21.06N, 55.73E | *Maladera affinis* | MalSB\_31 | Y | Y |
| RS5420 | SB | -21.06N, 55.73E | *Maladera affinis* | MalSB\_32 | Y | Y |
| RS5421 | SB | -21.06N, 55.73E | *Maladera affinis* | MalSB\_45 | Y | Y |
| RS5422 | SB | -21.06N, 55.73E | *Maladera affinis* | MalSB\_46 | Y | Y |
| RS5423 | SB | -21.06N, 55.73E | *Maladera affinis* | MalSB\_47 | Y | Y |
| RS5424 | SB | -21.06N, 55.73E | *Maladera affinis* | MalSB\_48 | Y | Y |
| RSB068 | SB | -21.06N, 55.73E | *Aphodius* sp. | AphSB\_1 | Y | Y |
| RSB069 | SB | -21.06N, 55.73E | *Aphodius* sp. | AphSB\_1 | Y | Y |
| RSB070 | SB | -21.06N, 55.73E | *Aphodius* sp. | AphSB\_1 | Y | Y |
| RSB071 | SB | -21.06N, 55.73E | *Aphodius* sp. | AphSB\_1 | Y | Y |
| RSB072 | SB | -21.06N, 55.73E | *Aphodius* sp. | AphSB\_2 | Y | Y |
| RSB073 | SB | -21.06N, 55.73E | *Aphodius* sp. | AphSB\_2 | Y | Y |
| RSB074 | SB | -21.06N, 55.73E | *Aphodius* sp. | AphSB\_2 | Y | Y |
| RSB075 | SB | -21.06N, 55.73E | *Aphodius* sp. | AphSB\_2 | Y | Y |
| RSB077 | SB | -21.06N, 55.73E | *Aphodius* sp. | AphSB\_8 | Y | Y |
| RSB078 | SB | -21.06N, 55.73E | *Aphodius* sp. | AphSB\_8 | Y | Y |
| RSB079 | SB | -21.06N, 55.73E | *Aphodius* sp. | AphSB\_8 | Y | Y |
| RSB080 | SB | -21.06N, 55.73E | *Aphodius* sp. | AphSB\_8 | Y | Y |
| RSB081 | SB | -21.06N, 55.73E | *Maladera affinis* | AphSB\_8 | Y | Y |
| RSB082 | SB | -21.06N, 55.73E | *Maladera affinis* | MalSB\_1 | Y | Y |
| RSB083 | SB | -21.06N, 55.73E | *Maladera affinis* | MalSB\_1 | Y | Y |
| RSB084 | SB | -21.06N, 55.73E | *Maladera affinis* | MalSB\_1 | Y | Y |
| RSB085 | SB | -21.06N, 55.73E | *Maladera affinis* | MalSB\_1 | Y | Y |
| RSB086 | SB | -21.06N, 55.73E | *Aphodius* sp. | MalSB\_1 | Y | Y |
| RSA096 | SS | -21.02N, 55.37E | *Oryctes borbonicus* | OrySS\_1 | Y | Y |
| RSA097 | SS | -21.02N, 55.37E | *Oryctes borbonicus* | OrySS\_1 | Y | Y |
| RSA098 | SS | -21.02N, 55.37E | *Oryctes borbonicus* | OrySS\_2 |  | Y |
| RSA100 | SS | -21.02N, 55.37E | *Oryctes borbonicus* | OrySS\_3 |  | Y |
| RSA101 | SS | -21.02N, 55.37E | *Oryctes borbonicus* | OrySS\_3 | Y | Y |
| RSA102 | SS | -21.02N, 55.37E | *Oryctes borbonicus* | OrySS\_7 | Y | Y |
| RSA103 | SS | -21.02N, 55.37E | *Oryctes borbonicus* | OrySS\_8 | Y | Y |
| RSA104 | SS | -21.02N, 55.37E | *Oryctes borbonicus* | OrySS\_8 |  | Y |
| RSA105 | SS | -21.02N, 55.37E | *Oryctes borbonicus* | OrySS\_9 | Y | Y |
| RSA106 | SS | -21.02N, 55.37E | *Oryctes borbonicus* | OrySS\_9 | Y | Y |
| RSA107 | SS | -21.02N, 55.37E | *Oryctes borbonicus* | OrySS\_9 |  | Y |
| RSA108 | SS | -21.02N, 55.37E | *Oryctes borbonicus* | OrySS\_10 | Y | Y |
| RSA109 | SS | -21.02N, 55.37E | *Oryctes borbonicus* | OrySS\_10 |  | Y |
| RSA110 | SS | -21.02N, 55.37E | *Oryctes borbonicus* | OrySS\_10 | Y | Y |
| RSA111 | SS | -21.02N, 55.37E | *Oryctes borbonicus* | OrySS\_12 | Y | Y |
| RS5333 | TB | -21.10N, 55.34E | *Oryctes borbonicus* | OryTB\_4 | Y | Y |
| RS5334 | TB | -21.10N, 55.34E | *Oryctes borbonicus* | OryTB\_8 | Y | Y |
| RS5336 | TB | -21.10N, 55.34E | *Oryctes borbonicus* | OryTB\_9 | Y | Y |
| RS5337 | TB | -21.10N, 55.34E | *Oryctes borbonicus* | OryTB\_10 | Y | Y |
| RS5338 | TB | -21.10N, 55.34E | *Oryctes borbonicus* | OryTB\_11 |  | Y |
| RS5339 | TB | -21.10N, 55.34E | *Oryctes borbonicus* | OryTB\_12 |  | Y |
| RS5346 | TB | -21.10N, 55.34E | *Oryctes borbonicus* | OryTB\_1 |  | Y |
| RS5347 | TB | -21.10N, 55.34E | *Oryctes borbonicus* | OryTB\_1 | Y | Y |
| RS5348 | TB | -21.10N, 55.34E | *Oryctes borbonicus* | OryTB\_1 |  | Y |
| RS5349 | TB | -21.10N, 55.34E | *Hoplia retusa* | HopTB\_1 |  | Y |
| RS5351 | TB | -21.10N, 55.34E | *Oryctes borbonicus* | OryTB\_2 | Y | Y |
| RS5353 | TB | -21.10N, 55.34E | *Oryctes borbonicus* | OryTB\_5 |  | Y |
| RS5354 | TB | -21.10N, 55.34E | *Oryctes borbonicus* | OryTB\_5 |  | Y |
| RS5355 | TB | -21.10N, 55.34E | *Oryctes borbonicus* | OryTB\_8 |  | Y |
| RS5377 | TB | -21.10N, 55.34E | *Oryctes borbonicus* | OryTB\_1 |  | Y |
| RS5378 | TB | -21.10N, 55.34E | *Oryctes borbonicus* | OryTB\_2 | Y | Y |
| RS5379 | TB | -21.10N, 55.34E | *Oryctes borbonicus* | OryTB\_2 |  | Y |
| RS5380 | TB | -21.10N, 55.34E | *Oryctes borbonicus* | OryTB\_3 |  | Y |
| RS5382 | TB | -21.10N, 55.34E | *Oryctes borbonicus* | OryTB\_3 |  | Y |
| RS5383 | TB | -21.10N, 55.34E | *Oryctes borbonicus* | OryTB\_3 |  | Y |
| RS5384 | TB | -21.10N, 55.34E | *Oryctes borbonicus* | OryTB\_4 | Y | Y |
| RS5385 | TB | -21.10N, 55.34E | *Oryctes borbonicus* | OryTB\_4 | Y | Y |
| RS5386 | TB | -21.10N, 55.34E | *Oryctes borbonicus* | OryTB\_5 |  | Y |
| RS5387 | TB | -21.10N, 55.34E | *Oryctes borbonicus* | OryTB\_6 |  | Y |
| RS5388 | TB | -21.10N, 55.34E | *Oryctes borbonicus* | OryTB\_6 |  | Y |
| RS5389 | TB | -21.10N, 55.34E | *Oryctes borbonicus* | OryTB\_8 |  | Y |
| RS5390 | TB | -21.10N, 55.34E | *Oryctes borbonicus* | OryTB\_9 |  | Y |
| RS5391 | TB | -21.10N, 55.34E | *Oryctes borbonicus* | OryTB\_9 |  | Y |
| RS5392 | TB | -21.10N, 55.34E | *Oryctes borbonicus* | OryTB\_10 |  | Y |
| RS5393 | TB | -21.10N, 55.34E | *Oryctes borbonicus* | OryTB\_10 |  | Y |
| RS5394 | TB | -21.10N, 55.34E | *Oryctes borbonicus* | OryTB\_10 | Y | Y |
| RS5395 | TB | -21.10N, 55.34E | *Oryctes borbonicus* | OryTB\_11 |  | Y |
| RS5396 | TB | -21.10N, 55.34E | *Oryctes borbonicus* | OryTB\_11 |  | Y |
| RS5397 | TB | -21.10N, 55.34E | *Oryctes borbonicus* | OryTB\_12 | Y | Y |
| RS5398 | TB | -21.10N, 55.34E | *Oryctes borbonicus* | OryTB\_12 | Y | Y |
| RS5399 | TB | -21.10N, 55.34E | *Oryctes borbonicus* | OryTB\_16 | Y | Y |
| RS5400 | TB | -21.10N, 55.34E | *Oryctes borbonicus* | OryTB\_16 | Y | Y |
| RS5401 | TB | -21.10N, 55.34E | *Oryctes borbonicus* | OryTB\_16 | Y | Y |
| RS5402 | TB | -21.10N, 55.34E | *Hoplia retusa* | HopTB\_1 | Y | Y |
| RS5403 | TB | -21.10N, 55.34E | *Hoplia retusa* | HopTB\_2 | Y | Y |
| RS5404 | TB | -21.10N, 55.34E | *Hoplia retusa* | HopTB\_7 |  | Y |
| RS5405 | TB | -21.10N, 55.34E | *Hoplia retusa* | HopTB\_15 | Y | Y |
| RSA001 | TB | -21.10N, 55.34E | *Oryctes borbonicus* | OryTB\_5 | Y | Y |
| RSA002 | TB | -21.10N, 55.34E | *Oryctes borbonicus* | OryTB\_5 | Y |  |
| RSA003 | TB | -21.10N, 55.34E | *Oryctes borbonicus* | OryTB\_5 | Y |  |
| RSA004 | TB | -21.10N, 55.34E | *Oryctes borbonicus* | OryTB\_5 | Y | Y |
| RSA005 | TB | -21.10N, 55.34E | *Oryctes borbonicus* | OryTB\_5 | Y |  |
| RSA006 | TB | -21.10N, 55.34E | *Oryctes borbonicus* | OryTB\_5 | Y |  |
| RSA007 | TB | -21.10N, 55.34E | *Oryctes borbonicus* | OryTB\_5 | Y |  |
| RSA008 | TB | -21.10N, 55.34E | *Oryctes borbonicus* | OryTB\_5 | Y |  |
| RSA009 | TB | -21.10N, 55.34E | *Oryctes borbonicus* | OryTB\_5 | Y |  |
| RSA010 | TB | -21.10N, 55.34E | *Oryctes borbonicus* | OryTB\_7 | Y |  |
| RSA011 | TB | -21.10N, 55.34E | *Oryctes borbonicus* | OryTB\_7 | Y | Y |
| RSA012 | TB | -21.10N, 55.34E | *Oryctes borbonicus* | OryTB\_7 | Y |  |
| RSA013 | TB | -21.10N, 55.34E | *Oryctes borbonicus* | OryTB\_7 | Y |  |
| RSA014 | TB | -21.10N, 55.34E | *Oryctes borbonicus* | OryTB\_7 | Y |  |
| RSA015 | TB | -21.10N, 55.34E | *Oryctes borbonicus* | OryTB\_7 | Y |  |
| RSA016 | TB | -21.10N, 55.34E | *Oryctes borbonicus* | OryTB\_7 | Y |  |
| RSA017 | TB | -21.10N, 55.34E | *Oryctes borbonicus* | OryTB\_15 | Y | Y |
| RSA018 | TB | -21.10N, 55.34E | *Oryctes borbonicus* | OryTB\_15 | Y | Y |
| RSA019 | TB | -21.10N, 55.34E | *Oryctes borbonicus* | OryTB\_15 | Y | Y |
| RSA020 | TB | -21.10N, 55.34E | *Oryctes borbonicus* | OryTB\_15 | Y |  |
| RSA021 | TB | -21.10N, 55.34E | *Oryctes borbonicus* | OryTB\_15 | Y |  |
| RSA022 | TB | -21.10N, 55.34E | *Oryctes borbonicus* | OryTB\_15 | Y |  |
| RSA023 | TB | -21.10N, 55.34E | *Oryctes borbonicus* | OryTB\_15 | Y |  |
| RSA027 | TB | -21.10N, 55.34E | *Oryctes borbonicus* | OryTB\_5 | Y |  |
| RSA028 | TB | -21.10N, 55.34E | *Oryctes borbonicus* | OryTB\_15 | Y |  |
| RSA029 | TB | -21.10N, 55.34E | *Oryctes borbonicus* | OryTB\_15 | Y |  |
| RSA030 | TB | -21.10N, 55.34E | *Oryctes borbonicus* | OryTB\_15 | Y | Y |
| RSA031 | TB | -21.10N, 55.34E | *Oryctes borbonicus* | OryTB\_15 | Y |  |
| RSA062 | TB | -21.10N, 55.34E | *Hoplochelus sp.* | HchTB\_5 |  | Y |
| RSA063 | TB | -21.10N, 55.34E | *Hoplochelus sp.* | HchTB\_5 | Y | Y |
| RSA064 | TB | -21.10N, 55.34E | *Hoplochelus sp.* | HchTB\_6 | Y | Y |
| RSA065 | TB | -21.10N, 55.34E | *Hoplochelus sp.* | HchTB\_6 | Y | Y |
| RSA066 | TB | -21.10N, 55.34E | *Hoplochelus sp.* | HchTB\_7 |  | Y |
| RSA067 | TB | -21.10N, 55.34E | *Hoplochelus sp.* | HchTB\_7 | Y | Y |
| RSA068 | TB | -21.10N, 55.34E | *Hoplochelus sp.* | HchTB\_9 |  | Y |
| RSA069 | TB | -21.10N, 55.34E | *Hoplochelus sp.* | HchTB\_9 | Y | Y |
| RSA070 | TB | -21.10N, 55.34E | *Hoplochelus sp.* | HchTB\_10 |  | Y |
| RSA071 | TB | -21.10N, 55.34E | *Hoplochelus sp.* | HchTB\_10 | Y | Y |
| RSA072 | TB | -21.10N, 55.34E | *Hoplochelus sp.* | HchTB\_11 | Y | Y |
| RSA073 | TB | -21.10N, 55.34E | *Hoplochelus sp.* | HchTB\_13 | Y | Y |
| RSA074 | TB | -21.10N, 55.34E | *Hoplochelus sp.* | HchTB\_13 |  | Y |
| RSA089 | TB | -21.10N, 55.34E | *Hoplia retusa* | HopTB\_4 | Y | Y |
| RSA090 | TB | -21.10N, 55.34E | *Hoplia retusa* | HopTB\_5 | Y | Y |
| RSA091 | TB | -21.10N, 55.34E | *Hoplia retusa* | HopTB\_6 | Y | Y |
| RSA092 | TB | -21.10N, 55.34E | *Hoplia retusa* | HopTB\_7 | Y | Y |
| RSA112 | TB | -21.10N, 55.34E | *Oryctes borbonicus* | OryTB\_8 |  | Y |
| RSA113 | TB | -21.10N, 55.34E | *Oryctes borbonicus* | OryTB\_17 | Y | Y |
| RSA114 | TB | -21.10N, 55.34E | *Oryctes borbonicus* | OryTB\_19 | Y | Y |
| RSB088 | TB | -21.10N, 55.34E | *Hoplochelus sp.* | HchTB\_2 | Y |  |
| RSB089 | TB | -21.10N, 55.34E | *Hoplochelus sp.* | HchTB\_2 | Y | Y |
| RSB090 | TB | -21.10N, 55.34E | *Hoplochelus sp.* | HchTB\_2 | Y | Y |
| RSB091 | TB | -21.10N, 55.34E | *Hoplochelus sp.* | HchTB\_2 | Y | Y |
| RSB092 | TB | -21.10N, 55.34E | *Hoplochelus sp.* | HchTB\_2 | Y | Y |
| RSB093 | TB | -21.10N, 55.34E | *Hoplia retusa* | HopTB\_4 | Y | Y |
| RSB094 | TB | -21.10N, 55.34E | *Hoplia retusa* | HopTB\_4 | Y | Y |
| RSB095 | TB | -21.10N, 55.34E | *Hoplia retusa* | HopTB\_4 | Y | Y |
| RSB096 | TB | -21.10N, 55.34E | *Hoplia retusa* | HopTB\_4 | Y | Y |
| RS5425 | TBG | -21.06N, 55.18E | *Aphodius sp.* | AphTBG\_2 |  | Y |
| RS5426 | TBG | -21.06N, 55.18E | *Hoplochelus sp.* | HchTBG\_2 |  | Y |
| RS5427 | TBG | -21.06N, 55.18E | *Hoplochelus sp.* | HchTBG\_2 | Y | Y |
| RS5428 | TBG | -21.06N, 55.18E | *Hoplochelus sp.* | HchTBG\_6 | Y | Y |
| RS5429 | TBG | -21.06N, 55.18E | *Hoplochelus sp.* | HchTBG\_7 | Y | Y |
| RS5430 | TBG | -21.06N, 55.18E | *Maladera affinis* | MalTBG\_1 | Y | Y |
| RSA060 | TBG | -21.06N, 55.18E | *Aphodius sp.* | AphTBG\_2 |  | Y |
| RSA093 | TBG | -21.06N, 55.18E | *Hoplia retusa* | HopTBG\_5 | Y | Y |
| RSA115 | TBG | -21.06N, 55.18E | *Oryctes borbonicus* | OryTBG\_1 |  | Y |
| RSA116 | TBG | -21.06N, 55.18E | *Oryctes borbonicus* | OryTBG\_2 |  | Y |
| RSA117 | TBG | -21.06N, 55.18E | *Oryctes borbonicus* | OryTBG\_2 | Y | Y |
| RSB127 | TBG | -21.06N, 55.18E | *Ataenius* sp. | AtaTBG\_21:29 |  | Y |
| RSB128 | TBG | -21.06N, 55.18E | *Ataenius* sp. | AtaTBG\_21:29 |  | Y |
| RSB130 | TBG | -21.06N, 55.18E | *Ataenius* sp. | AtaTBG\_21:29 |  | Y |
| RSB131 | TBG | -21.06N, 55.18E | *Ataenius* sp. | AtaTBG\_21:29 |  | Y |
| RSB132 | TBG | -21.06N, 55.18E | *Ataenius* sp. | AtaTBG\_21:29 |  | Y |
| RSB133 | TBG | -21.06N, 55.18E | *Ataenius* sp. | AtaTBG\_21:29 |  | Y |
| RSB134 | TBG | -21.06N, 55.18E | *Ataenius* sp. | AtaTBG\_21:29 |  | Y |
| RSB135 | TBG | -21.06N, 55.18E | *Hoplochelus sp.* | HchTBG\_1 |  | Y |
| RSB136 | TBG | -21.06N, 55.18E | *Hoplochelus sp.* | HchTBG\_1 |  | Y |
| RSB138 | TBG | -21.06N, 55.18E | *Hoplochelus sp.* | HchTBG\_1 |  | Y |
| RSB139 | TBG | -21.06N, 55.18E | *Hoplochelus sp.* | HchTBG\_1 |  | Y |
| RSB142 | TBG | -21.06N, 55.18E | *Hoplochelus sp.* | HchTBG\_4 |  | Y |
| RSB143 | TBG | -21.06N, 55.18E | *Hoplochelus sp.* | HchTBG\_4 |  | Y |
| RSB144 | TBG | -21.06N, 55.18E | *Hoplochelus sp.* | HchTBG\_4 |  | Y |
| RSB145 | TBG | -21.06N, 55.18E | *Hoplochelus sp.* | HchTBG\_4 |  | Y |