|  |  |  |  |
| --- | --- | --- | --- |
| **Supplementary file 5. DNA sequencing yield.** | | | |
| **Sample ID** | **Sampling location** | **Ecotype** | **Mapped reads \*** |
| FC14 | Little Campbell River | Freshwater | 17438204 |
| FC13 | Little Campbell River | Freshwater | 23171229 |
| FC12 | Little Campbell River | Freshwater | 25146843 |
| FC16 | Little Campbell River | Freshwater | 30005091 |
| FC18 | Little Campbell River | Freshwater | 38399784 |
| FC15 | Little Campbell River | Freshwater | 15152985 |
| c363\_P\_FC08\_F | Little Campbell River | Freshwater (parent) | 58346540 |
| LITC\_DWN\_4 | Little Campbell River | Marine | 28238410 |
| LITC\_DWN\_5\_F | Little Campbell River | Marine | 37997779 |
| LITC\_DWN\_6\_F | Little Campbell River | Marine | 46675891 |
| LITC\_DWN\_7\_F | Little Campbell River | Marine | 27626027 |
| LITC\_DWN\_8\_F | Little Campbell River | Marine | 30613683 |
| LITC\_DWN\_9\_F | Little Campbell River | Marine | 25885479 |
| c363\_P\_FC18\_M | Little Campbell River | Marine (parent) | 36326200 |
| Tyne8\_27 | River Tyne | Freshwater | 57821343 |
| Tyne8\_4 | River Tyne | Freshwater | 75419343 |
| Tyne8\_7 | River Tyne | Freshwater | 80784624 |
| Tank422\_4 | River Tyne | Freshwater | 120054557 |
| Tyne8\_2 | River Tyne | Freshwater | 89618061 |
| Tyne8\_1 | River Tyne | Freshwater | 58244138 |
| c172\_P\_533\_M | River Tyne | Freshwater (parent) | 56617456 |
| Tyne2\_16|2015 | River Tyne | Marine | 89914271 |
| Tyne2\_18 | River Tyne | Marine | 28775921 |
| Tyne2\_20 | River Tyne | Marine | 55538908 |
| Tyne2\_12 | River Tyne | Marine | 15830364 |
| Tyne2\_16|2014 | River Tyne | Marine | 24148885 |
| Tyne2\_14 | River Tyne | Marine | 108126294 |
| c172\_P\_532\_F | River Tyne | Marine (parent) | 66347185 |
| c214\_P\_512\_M | River Shiel | Freshwater (parent) | 57188122 |
| c214\_P\_524\_F | River Shiel | Marine (parent) | 61037784 |
| c212\_P\_551\_M | River Forss | Freshwater (parent) | 53922480 |
| c212\_P\_454\_F | River Forss | Marine (parent) | 27969046 |
| \* samtools view -F 0x4 \*.bam | cut -f 1 | sort | uniq | wc -l | | | |