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| **Table S1.** *Hypomesus nipponensis* genome sequencing. | | |
| **Sequencing type** | **Yields** | **Length of reads** |
| Sequel (>20Kb) | 58.60Gbp | 5.7M |
| Novaseq6000 (2x101bp) | 36.99Gbp | 366.2M |