

Rate [%]	Mask	Hypothesis
19,03	-genome-	Unligated genome
6,98	-genome--a-bridge-gatc--egdirb-t--genome-	Expected ligation product
5,92	-genome--bridge--gatc-egdirb-t--genome-	Unexpected ligation product
5,83	-genome--a-bridge-gatc--egdirb--genome-	Unexpected ligation product
5,62	-genome--bridge-gatc--egdirb--genome-	Unexpected ligation product
4,59	-genome--a-bridge-gatc--genome-	Unexpected ligation product
4,05	-genome--bridge-gatc--genome-	Unexpected ligation product
3,14	-genome--gatc-egdirb-t--genome-	Unexpected ligation product
2,30	-genome--gatc-egdirb--genome-	Unexpected ligation product
0,99	-genome--bridge--genome-	Unexpected ligation product
0,99	-genome--egdirb--genome-	Unexpected ligation product
0,93	-genome--egdirb-t--genome-	Unexpected ligation product
0,72	-genome--a-bridge-gatc--egdirb-	Unexpected ligation product
0,71	-genome--a-bridge--genome-	Unexpected ligation product
0,64	-bridge--gatc-egdirb-t--genome-	Unexpected ligation product
0,59	--[6]--gatc-egdirb-t--genome-	Unexpected ligation product
0,54	-bridge-gatc--egdirb--genome-	Unexpected ligation product
0,51	-bridge-gatc--genome-	Unexpected ligation product
0,49	-genome--a-bridge-gatc--[6]--	Unexpected ligation product
0,47	-genome--bridge-gatc--egdirb-	Unexpected ligation product