

Rate [%]	Mask	Hypothesis
16,52	-genome-	Unligated genome
7,86	-genome--bridge-gatc--egdirb--genome-	Unexpected ligation product
6,83	-genome--a-bridge-gatc--egdirb--genome-	Unexpected ligation product
6,41	-genome--bridge--gatc-egdirb-t--genome-	Unexpected ligation product
3,39	-genome--a-bridge-gatc--egdirb-t--genome-	Expected ligation product
2,28	-genome--bridge-gatc--egdirb--bridge-	Unexpected ligation product
2,26	-egdirb--bridge-gatc--egdirb--genome-	Unexpected ligation product
2,24	-genome--gatc-egdirb--genome-	Unexpected ligation product
2,12	-genome--gatc-egdirb-	Unexpected ligation product
1,97	-bridge-gatc--genome-	Unexpected ligation product
1,91	-genome--gatc-egdirb--bridge-	Unexpected ligation product
1,90	-genome--bridge-gatc--genome-	Unexpected ligation product
1,88	-genome--a-bridge-gatc--egdirb--bridge-	Unexpected ligation product, adapter multimers
1,83	-egdirb--bridge--gatc-egdirb-t--genome-	Unexpected ligation product, adapter multimers
1,47	-genome--egdirb--genome-	Unexpected ligation product
1,41	-genome--gatc-egdirb-t--genome-	Unexpected ligation product
1,40	-egdirb--bridge-gatc--genome-	Unexpected ligation product
1,29	-genome--bridge--genome-	Unexpected ligation product
1,27	-genome--a-bridge-gatc--genome-	Unexpected ligation product
0,97	-genome--bridge-gatc--egdirb-	Unexpected ligation product