Convolutio	Encoder	ecoder	Evaluated	Failed aligr	Short align	o predicti	Match	Mismatch	nsertion	Deletion	Adenine	Cytosine	Guanine	Thymine	AUC	PhredQ ov
	Д	Dé l'	— — — — — — — — — — — — — — — — — — —	- Fa	No.	- N	Σ̈́	Σ	<u> </u>		- Ac		- B	<u>-</u>	-AL	- L
URN	LS5	CRF	99.4	0.20	0.32	0.06	93.3	2.08	1.86	2.76	9.94	13.1	16.4	11.9	94.4	20.6
<mark>URN</mark>	BFW	CRF	99.4	0.21	0.33	0.06	93.3	2.04	1.72	2.89	9.62	13.5	16.3	11.9	94.4	19.0
CAT	BFW	CRF	99.4	0.23	0.33	0.05	93.4	1.98	1.56	3.00	9.90	14.2	16.6	12.2	94.5	18.1
CAT	LS5	CRF	99.4	0.25	0.32	0.05	93.6	1.94	1.47	3.02	10.0	14.0	16.5	12.5	94.5	21.0
BON	LS5	CRF	99.4	0.25	0.32	0.01	93.3	2.03	1.74	2.92	9.65	14.2	17.1	12.5	94.3	22.3
CAU	LS5	CRF	99.4	0.23	0.33	0.04	93.2	2.11	1.88	2.79	9.15	14.0	17.0	11.5	94.3	21.2
BON	BRE	CRF	99.4	0.25	0.32	0.02	93.0	2.16	1.79	3.07	9.73	16.0	17.7	13.8	93.9	24.5
SAC	LS5	CRF	99.4	0.19	0.33	0.09	93.2	2.07	1.61	3.10	10.6	15.0	19.1	14.0	94.2	19.0
BON	BFW	CRF	99.4	0.22	0.33	0.02	92.8	2.23	1.97	2.98	10.0	15.1	19.2	13.6	93.8	17.9
URN,	LS3	CRF	99.4	0.21	0.32	0.08	92.8	2.22	1.92	3.02	9.99	15.3	18.8	13.7	93.7	21.9

⊤ Better

-Company Derformance change

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