

Protein	# of positives	precision	recall	f1-score	roc-auc	pr-auc
AGO1	31964	0.862	0.747	0.801	0.947	0.901
LIN28A	12821	0.898	0.721	0.8	0.946	0.862
TIAL1	25616	0.836	0.718	0.772	0.936	0.869
TIA1	19453	0.807	0.733	0.768	0.946	0.853
LIN28B	15063	0.901	0.67	0.768	0.932	0.84
RBM47	18653	0.83	0.657	0.733	0.946	0.838
LARP4B	18832	0.756	0.662	0.706	0.92	0.79
AGO2	20761	0.771	0.609	0.681	0.909	0.767
FMR1_iso1	16923	0.871	0.558	0.68	0.923	0.803
Negatives	31964	0.728	0.636	0.679	0.87	0.766
ZC3H7B	12439	0.801	0.581	0.674	0.939	0.769
ELAVL1	25715	0.818	0.567	0.67	0.879	0.775
ATXN2	10425	0.736	0.583	0.651	0.93	0.73
FXR2	10460	0.877	0.515	0.649	0.924	0.751
MOV10	6980	0.766	0.543	0.635	0.956	0.731
TAF15	4462	0.817	0.504	0.623	0.941	0.669
EIF3A	559	0.828	0.495	0.62	0.967	0.678
FXR1	3358	0.847	0.486	0.617	0.959	0.719
EWSR1	6214	0.765	0.508	0.611	0.951	0.698
SRRM4	25381	0.721	0.519	0.604	0.86	0.715
FMR1_iso7	18228	0.823	0.476	0.603	0.897	0.737
PCBP2	3749	0.732	0.493	0.589	0.926	0.627
YTHDF1	6648	0.792	0.455	0.578	0.927	0.673
CAPRIN1	7012	0.703	0.472	0.565	0.919	0.631
MSI1	10801	0.741	0.454	0.563	0.904	0.646
FUS	7577	0.741	0.451	0.561	0.923	0.644
EIF4A3	21759	0.719	0.447	0.552	0.869	0.677
HNRNPC	4799	0.726	0.417	0.529	0.948	0.598
EIF3D	615	0.852	0.351	0.497	0.963	0.65
EIF3B	421	0.725	0.362	0.483	0.964	0.566
C22ORF28	5938	0.747	0.353	0.479	0.889	0.538
YTHDF2	2108	0.936	0.294	0.447	0.924	0.629
HNRNPD	15786	0.739	0.303	0.43	0.817	0.557
HNRNPH1	4858	0.876	0.284	0.429	0.926	0.604
EIF3G	597	0.841	0.278	0.418	0.97	0.67
RBM10	9968	0.72	0.287	0.41	0.869	0.534
IGF2BP2	9265	0.534	0.325	0.404	0.858	0.431
HNRNPF	4503	0.778	0.269	0.4	0.898	0.497
IGF2BP1	9389	0.531	0.316	0.396	0.849	0.424
IGF2BP3	11429	0.574	0.28	0.376	0.849	0.443
DDX21	9424	0.691	0.256	0.373	0.857	0.487
ZFP36	4323	0.766	0.245	0.372	0.936	0.565
RBPMS	4706	0.655	0.214	0.322	0.929	0.472
ADAR1	2210	0.83	0.173	0.286	0.928	0.427
HNRNPU	4736	0.644	0.146	0.238	0.89	0.392
PUM2	3581	0.515	0.148	0.23	0.906	0.303

TARDBP	1332	0.806	0.109	0.191	0.926	0.341
TARBP2	460	0.8	0.094	0.168	0.974	0.428
CPEB1	716	0.818	0.077	0.141	0.884	0.252
RBFOX2	850	0.632	0.073	0.131	0.891	0.224
HNRNPA2B1	2201	0.743	0.061	0.113	0.899	0.285
WDR33	902	0.833	0.054	0.102	0.94	0.382
ALKBH5	1165	0.667	0.055	0.102	0.907	0.29
PUM1	3788	0.767	0.043	0.081	0.775	0.191
ZC3H12A	799	1	0.041	0.078	0.932	0.297
METTL3	729	0.833	0.039	0.075	0.891	0.215
CELF1	940	1	0.034	0.065	0.88	0.194
STAU1	3520	0.875	0.029	0.057	0.776	0.202
TNRC6B	1070	0.286	0.029	0.053	0.863	0.093
DGCR8	1600	1	0.023	0.044	0.863	0.197
HNRNPA1	632	0.5	0.016	0.031	0.906	0.242
PARK7	697	1	0.015	0.029	0.804	0.093
PABPC1	2322	1	0.008	0.017	0.754	0.124
C17ORF85	1386	0.5	0.006	0.012	0.848	0.183
QKI	1008	0.5	0.005	0.01	0.867	0.124
U2AF2	2202	1	0.003	0.005	0.728	0.101
RC3H1	2950	1	0.002	0.003	0.712	0.09
AGO4	400	0	0	0	0.882	0.043