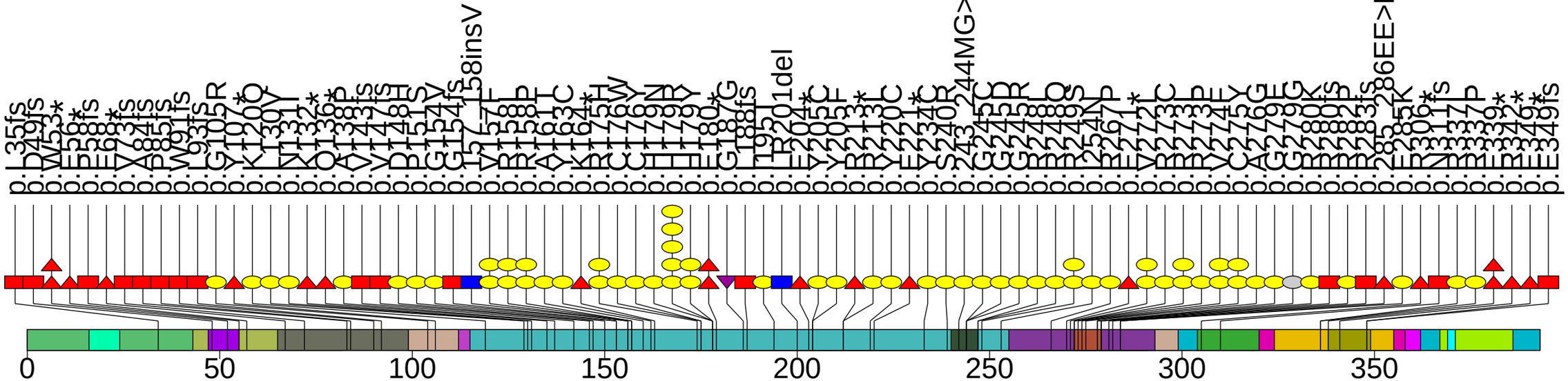
p.D19fs
p.W53*
p.L56*fs
p.E69*fs
p.V72fs
p.A85fs
p.W91fs
p.G105R
p.Y107*
p.K120Q
p.L130Y
p.K132*
p.Q136*
p.L138fs
p.Y147fs
p.D148H
p.R152S
p.G154V
p.G154fs
p.L157F
p.Y158F
p.R158P
p.R161L
p.Q163C
p.K164*
p.R175H
p.C176W
p.C176Y
p.T178N
p.T178R
p.T179P
p.T180*
p.G187G
p.T188fs
p.L195I
p.R201del
p.E204*
p.Y205C
p.Y205F
p.R213*
p.R213L
p.E220C
p.E221*
p.Y234C
p.S240R
p.243_244MG>L
p.G245C
p.G245D
p.G245R
p.R248Q
p.R248W
p.R249S
p.R249M
p.R250L
p.E267*
p.V272L
p.R273C
p.R273H
p.R273P
p.R273L
p.V274F
p.C275Y
p.A276G
p.G279E
p.G279K
p.R280*fs
p.R282P
p.R283fs
p.286_285EE>K
p.E285K
p.R306*
p.N307fs
p.R337L
p.R337P
p.R339*
p.E342*
p.E346*
p.E349fs
0
50
100
150
200
250
300
350
DNA−binding (experimentally determined)
Interaction with CCAR2. {ECO:0000269|PubMed:257328
Interaction with HRMT1L2. {ECO:0000269|PubMed:1518
Transcription activation (acidic).
Interaction with WWOX.
Interaction with HIPK1. ECO:0000250.
Required for interaction with ZNF385A. {ECO:000026
Required for interaction with FBXO42. {ECO:0000269
Interaction with AXIN1. ECO:0000250.
Interaction with the 53BP2 SH3 domain.
Interaction with E4F1. {ECO:0000269|PubMed:1064499
Interaction with DNA.
Interaction with CARM1. {ECO:0000269|PubMed:151867
Interaction with HIPK2.
Oligomerization.
Interaction with USP7.
Basic (repression of DNA−binding).
TADI.
TADII.
Bipartite nuclear localization signal.
Nuclear export signal.
[KR]−[STA]−K motif.
Silent
In_Frame_Del
Splice_Site
In_Frame_Ins
Missense_Mutation
Frame_Shift_Ins
Nonsense_Mutation
Frame_Shift_Del