

miRNAfe: multidimensional features name description

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dinucleotide_proportion0	AA
dinucleotide_proportion1	AC
dinucleotide_proportion2	AG
dinucleotide_proportion3	AU
dinucleotide_proportion4	CA
dinucleotide_proportion5	CC
dinucleotide_proportion6	CG
dinucleotide_proportion7	CU
dinucleotide_proportion8	GA
dinucleotide_proportion9	GC
dinucleotide_proportion10	GG
dinucleotide_proportion11	GU
dinucleotide_proportion12	UA
dinucleotide_proportion13	UC
dinucleotide_proportion14	UG
dinucleotide_proportion15	UU

huang_elements_proportion0	"="
huang_elements_proportion1	"=="
huang_elements_proportion2	"=:"
huang_elements_proportion3	"_:"
huang_elements_proportion4	"-="
huang_elements_proportion5	"^^"
huang_elements_proportion6	"^="
huang_elements_proportion7	"::"
huang_elements_proportion8	".^"
huang_elements_proportion9	".="

huang_ratios0	pMatch
huang_ratios1	pMismatch
huang_ratios2	pDI
huang_ratios3	pBulge

bp_proportion_stem0	A-U/Nstems
bp_proportion_stem1	G-C/Nstems
bp_proportion_stem2	G-U/Nstems

bp_proportion0	A-U
bp_proportion1	G-C
bp_proportion2	G-U

triplets0	((a
triplets1	((c
triplets2	((g
triplets3	((u
triplets4	((a
triplets5	((c
triplets6	((g
triplets7	((u
triplets8	((a
triplets9	((c
triplets10	((g
triplets11	((u
triplets12	.(a
triplets13	.(c
triplets14	.(g
triplets15	.(u
triplets16	.(a
triplets17	.(c
triplets18	.(g
triplets19	.(u
triplets20	.(a
triplets21	.(c
triplets22	.(g
triplets23	.(u
triplets24	..(a
triplets25	..(c
triplets26	..(g
triplets27	..(u
triplets28	...a
triplets29	...c
triplets30	...g
triplets31	...u

nt_porportion0	A
nt_porportion1	C
nt_porportion2	G
nt_porportion3	U