

Non-LTR retrotransposon の group と clade のリスト

これまでに提唱された non-LTR retrotransposon の group と clade をまとめてみました。最初に non-LTR retrotransposon の分類単位としての group と clade の定義を紹介します。

Clade は Malik らによって 1999 年に提唱された分類単位で、以下の三つの条件を満たすものとされました(Malik, Burke and Eickbush 1999)。

- (1) 構造上共通する特徴を持つ
- (2) 逆転写酵素を用いた系統解析で単系統性が十分支持される
- (3) 系統の起源が先カンブリア時代まで遡る

Malik らは 11 の clade を提唱しましたが、その後新しい clade が次々と提唱され、数が増加したため、Eickbush らにより group が提唱されました(Eickbush and Malik 2002)。Group は clade を更にまとめる単位で、逆転写酵素の系統樹とタンパク質のドメイン構造に基づいています。ここで提唱された五つの group (R2, L1, RTE, I, Jockey) に加えて、Dualen、CR1、CRE の 3 つが提唱されていますが、group はあくまで clade をまとめるために導入された便宜的な分類単位ですので、これ自体に科学的な意味はありません。

Table. Proposed Groups and Clades of non-LTR retrotransposons.

Group	Clade	Representative	Species	References
CRE	CRE	CRE2	<i>Crithidia fasciculata</i>	(Malik, Burke and Eickbush 1999)
	Genie	<i>Genie1A/GilM</i>	<i>Giardia lamblia</i>	(Burke et al. 2002)
R2	R2	<i>R2Dm</i>	<i>Drosophila melanogaster</i>	(Malik, Burke and Eickbush 1999)
	R2-A	<i>R2Dr</i>	<i>Danio rerio</i>	(Kojima and Fujiwara 2005b)
	R2-B	<i>R2-I_TCas</i>	<i>Tribolium castaneum</i>	(Kojima and Fujiwara 2005b)
	R2-C	<i>R2Sm-A</i>	<i>Schistosoma mansoni</i>	(Kojima and Fujiwara 2005b)
	R2-D	<i>R2Dm</i>	<i>Drosophila melanogaster</i>	(Kojima and Fujiwara 2005b)
R4	R4		<i>Ascaris lumbricoides</i>	(Malik, Burke and Eickbush 1999)

Non-LTR retrotransposon の group と clade のリスト

	NeSL	<i>NeSL-1</i>	<i>Caenorhabditis elegans</i>	(Malik and Eickbush 2000)
	HERO	<i>HERODr</i>	<i>Danio rerio</i>	(Kojima et al. 2006)
Dualen	Dualen	<i>DualenCrI</i>	<i>Chlamydomonas reinhardtii</i>	(Kojima and Fujiwara 2005a)
L1	L1	<i>L1</i>	<i>Homo sapiens</i>	(Malik, Burke and Eickbush 1999)
Tx1		<i>Tx1</i>	<i>Xenopus laevis</i>	(Putnam et al. 2007)
Proto1		<i>Proto1-1_NG</i>	<i>Naegleria gruberi</i>	(Kapitonov and Jurka 2009a)
RTE	RTE	<i>RTE-1</i>	<i>Caenorhabditis elegans</i>	(Malik, Burke and Eickbush 1999)
RTEX		<i>RTEX-1_NV</i>	<i>Nematostella vectensis</i>	(Putnam et al. 2007)
RTETP		<i>RTE-1_TP</i>	<i>Thalassiosira pseudonana</i>	(Kapitonov, Tempel and Jurka 2009)
Proto2		<i>Proto2-1_BF</i>	<i>Branchiostoma floridae</i>	(Kapitonov and Jurka 2009b)
I	I	<i>I</i>	<i>Drosophila melanogaster</i>	(Malik, Burke and Eickbush 1999)
Ingi		<i>Ingi</i>	<i>Trypanosoma brucei</i>	(Eickbush and Malik 2002)
Vingi		<i>Vingi-1_EE</i>	<i>Erinaceus europaeus</i>	(Kojima, Kapitonov and Jurka 2010)
Outcast		<i>Ag-Outcast-6</i>	<i>Anopheles gambiae</i>	(Biedler and Tu 2003)
Loner		<i>Ag-Loner-1</i>	<i>Anopheles gambiae</i>	(Biedler and Tu 2003)
Nimb		<i>nimbus</i>	<i>Biomphalaria glabrata</i>	(Kapitonov, Tempel and Jurka 2009)
R1		<i>RIBm</i>	<i>Bombyx mori</i>	(Malik, Burke and Eickbush 1999)
LOA		<i>LOA</i>	<i>Drosophila silvestris</i>	(Malik, Burke and Eickbush 1999)
Tad1		<i>Tad1</i>	<i>Neurospora crassa</i>	(Malik, Burke and Eickbush 1999)
Jockey	Jockey	<i>Jockey</i>	<i>Drosophila</i>	(Malik, Burke and Eickbush)

			<i>melanogaster</i>	1999)
CR1	CR1	<i>CRI</i>	<i>Gallus gallus</i>	(Malik, Burke and Eickbush 1999)
L2		<i>CRI-I_DR</i>	<i>Danio rerio</i>	(Lovsin, Gubensek and Kordi 2001)
Rex1		<i>REXI_DR</i>	<i>Danio rerio</i>	(Volff, Korting and Schartl 2000)
L2A		<i>CRI-34_HM</i>	<i>Hydra magnipapillata</i>	(Kapitonov, Tempel and Jurka 2009)
L2B		<i>CRI-I_AG</i>	<i>Anopheles gambiae</i>	(Kapitonov, Tempel and Jurka 2009)
Daphne		<i>Daphne_DS</i>	<i>Darwinula stevensoni</i> .	(Schon and Arkhipova 2006)
Crack				(Kapitonov, Tempel and Jurka 2009)
?	REP	<i>REPI</i>	<i>Tetrahymena thermophila</i>	(Fillingham et al. 2004)
?	Odin	<i>Odin-I</i>	<i>Oikopleura dioica</i>	(Volff et al. 2004)
?	Inkcap	<i>SrNLRI</i>	<i>Sporobolomyces roseus</i>	(Novikova, Fet and Blinov 2009)
?	Deceiver	<i>LbNLR8</i>	<i>Laccaria bicolor</i>	(Novikova, Fet and Blinov 2009)
?	Ambal	<i>Ambal-I_FCy</i>	<i>Fragilaropsis cylindrus</i>	(Kapitonov and Jurka 2010)

References

- Biedler J, Tu Z. 2003. Non-LTR retrotransposons in the African malaria mosquito, *Anopheles gambiae*: unprecedented diversity and evidence of recent activity. Mol Biol Evol 20:1811-1825.
- Burke WD, Malik HS, Rich SM, Eickbush TH. 2002. Ancient lineages of non-LTR retrotransposons in the primitive eukaryote, *Giardia lamblia*. Mol Biol Evol 19:619-630.
- Eickbush TH, Malik HS. 2002. Origins and evolution of retrotransposons. In: NL Craig, R Craigie, M Gellert, AM Lambowitz, editors. Mobile DNA II. Washington D. C.: American Society of

- Microbiology Press. p. 1111-1144.
- Fillingham JS, Thing TA, Vythilingum N, Keuroghlian A, Bruno D, Golding GB, Pearlman RE. 2004. A non-long terminal repeat retrotransposon family is restricted to the germ line micronucleus of the ciliated protozoan *Tetrahymena thermophila*. *Eukaryot Cell* 3:157-169.
- Kapitonov VV, Jurka J. 2009a. Proto1 non-LTR retrotransposons from the *Naegleria gruberi* amoeboflagellate genome. *Repbase Reports* 9:1144-1148.
- Kapitonov VV, Jurka J. 2009b. Proto2, a novel clade of metazoan non-LTR retrotransposons. *Repbase Reports* 9:1554-1563.
- Kapitonov VV, Jurka J. 2010. Ambal, a novel clade of non-LTR retrotransposons from diatoms. *Repbase Reports* 10:102-108.
- Kapitonov VV, Tempel S, Jurka J. 2009. Simple and fast classification of non-LTR retrotransposons based on phylogeny of their RT domain protein sequences. *Gene* 448:207-213.
- Kojima KK, Fujiwara H. 2005a. An extraordinary retrotransposon family encoding dual endonucleases. *Genome Res* 15:1106-1117.
- Kojima KK, Fujiwara H. 2005b. Long-term inheritance of the 28S rDNA-specific retrotransposon R2. *Mol Biol Evol* 22:2157-2165.
- Kojima KK, Kapitonov VV, Jurka J. 2010. Recent expansion of a new Ingi-related clade of Vingi non-LTR retrotransposons in hedgehogs. *Mol Biol Evol*.
- Kojima KK, Kuma K, Toh H, Fujiwara H. 2006. Identification of rDNA-specific non-LTR retrotransposons in Cnidaria. *Mol Biol Evol* 23:1984-1993.
- Lovsin N, Gubensek F, Kordi D. 2001. Evolutionary dynamics in a novel L2 clade of non-LTR retrotransposons in Deuterostomia. *Mol Biol Evol* 18:2213-2224.
- Malik HS, Burke WD, Eickbush TH. 1999. The age and evolution of non-LTR retrotransposable elements. *Mol Biol Evol* 16:793-805.
- Malik HS, Eickbush TH. 2000. NeSL-1, an ancient lineage of site-specific non-LTR retrotransposons from *Caenorhabditis elegans*. *Genetics* 154:193-203.
- Novikova O, Fet V, Blinov A. 2009. Non-LTR retrotransposons in fungi. *Funct Integr Genomics* 9:27-42.
- Putnam NH, Srivastava M, Hellsten U, et al. (XX co-authors). 2007. Sea anemone genome reveals ancestral eumetazoan gene repertoire and genomic organization. *Science* 317:86-94.
- Schon I, Arkhipova IR. 2006. Two families of non-LTR retrotransposons, Syrinx and Daphne, from the Darwinulid ostracod, *Darwinula stevensoni*. *Gene* 371:296-307.
- Volff JN, Korting C, Schartl M. 2000. Multiple lineages of the non-LTR retrotransposon Rex1 with varying success in invading fish genomes. *Mol Biol Evol* 17:1673-1684.

Non-LTR retrotransposon の group と clade のリスト

Volff JN, Lehrach H, Reinhardt R, Chourrout D. 2004. Retroelement dynamics and a novel type of chordate retrovirus-like element in the miniature genome of the tunicate *Oikopleura dioica*. Mol Biol Evol 21:2022-2033.

2010/12/13

小島 健司 著
禁 無断複写転載