

Table S1 Ozawa et al.

Comparison of the amino acid sequences between FLuc, CBR and ELuc.

1st Amino Acid Sequence File Name : PpyLuc Sequence Size : 550	1st Amino Acid Sequence File Name : PpyLuc Sequence Size : 550	1st Amino Acid Sequence File Name : ELuc Sequence Size : 542
2nd Amino Acid Sequence File Name : CBR Sequence Size : 542	2nd Amino Acid Sequence File Name : ELuc Sequence Size : 542	2nd Amino Acid Sequence File Name : CBR Sequence Size : 542
[48.8% / 543 aa] 1" MEDAKNIKKGPAPFPYPLEDTAGEQLHKAMKRYALVPGTIAFTDAHIEDVITYAEYFEM ...*, **, ****, ****, *, ..., *., *, ...*, ** 1" MVKREKNVIYGPPEPLHPLEDLTAGEMLFRALRKHSHL-P-AOLDVVGDESLYKEFFEA 60" SVRLAECAMKRYGLNTNHRIVVCSENSLSQFFMPVLGALFIGVAVAPANDIYNERELNSMG ...*, ..., *, **, **, **, **, **, **, **, **, ** 59" TVLLAQSLHNCGYKMNDVWSICAENTRRFIPVIAAWYIGMIVAPVNESYIPDELCKVMG 120" ISQPTVVFSKKLQKILNVQKQPIIQQIIIMDSKTDYQGFQSMTYFVTSHLPPGFNEY **., **, **, **, **, **, **, **, **, **, ** 119" ISKPQIVFTTKNILNKVLEVOVSRTNFIRIILDTVENIHGCESLPNF-SRYSDG-NIA 180" DFVPESFDRTKITALIMNSGSTGLPKGVALPHRTACVRFSHARDPIFGNQIPTDIALS **., **, **, **, **, **, **, **, **, **, **, ** 177" NFKPLHDFPVEQVAAILCSSLGTTLPKGVMQTHQNICVRILHALDPRTQQLIPGVTVL 240" VVPFHGFGMFITLGYLICGRVRLMYRFEELFLRSLODQYIQSALLVPTLSSFAKST ***, **, **, **, **, **, **, **, **, **, **, **, ** 237" YLPFFHAFGHITLGYFMVGLRVMFRRFDQEAFLKAIDQYEVRSINVPVSLFLSKSP 300" LIDKYDLNSLHEIASGGAPLSKEVEAVALKRFLPGLPQYQGLTETTSAILITPEGDDKP *****+, **, **, **, **, **, **, **, **, **, **, **, ** 297" LVDKYDLSSRELCGAAPLAKEVAEAAKRNLPGIRCGFLTESTSAIQTLDGFDFKS 360" GAVGVVPFFEAKWVLDLTGKTLGVNQRGECLVRGPIMISGYVNNPEATNALIDDGWLH ***, **, **, **, **, **, **, **, **, **, **, ** 357" GSLSRVLPLMAAKIADRETGKALGPQNQVGECLIKGPMVSKYVNNVTEAKIAIDDGGWL 420" SGDIAYWDEDEHFFIVDRLKSLIKYKGYQVAPAELESILQHPNIFDAGVAGLPDDDAEGE ***, **, **, **, **, **, **, **, **, **, **, ** 417" SGDFGYDDEDEHFYVDRYKELIKYKGQVAPAELEEILKNPCIRDVAVGIPDLEAGE 480" LPAAVVVLLEHGTMTKEIVDVAQSQTAAKKLRGGV/FVDEVPKGLTGKLDARK-IREI ***, **, **, **, **, **, **, **, **, **, ** 477" LPSAFVVKQPGTEITAKEVDYLAERVSHTKYLRRGVRFVDSIPRNVTGKTRKELLKQL 539" LIKAKGGKIAV ** 537" LVKAGG	[47.8% / 540 aa] 1" MEDAKNIKKGPAPFPYPLEDTAGEQLHKAMKRYALVPGTIAFTDAHIEDVITYAEYFEM ...*, **, **, **, ****, ****, *, ..., *., *, ...*, ** 1" MERENKNVYGPPEPKHPLGNFTAGEMLYNALHKHSHHQI-LDVMGNESLSYQEFFDTT 61" VRLAECAMKRYGLNTNHRIVVCSENSLSQFFMPVLGALFIGVAVAPANDIYNERELNSMG ...*, ..., *, **, **, **, **, **, **, **, ** 59" VKLGQSLQNCGYKMNDVWSICAENNRKFIPISAWSYIGMIVAPVNEDYIPDELCKVTG 121" SQPTVVFSKKLQKILNVQKQPIIQQIIIMDSKTDYQGFQSMTYFVTSHLPPGFNEYD **., **, **, **, **, **, **, **, **, **, ** 119" SKPILVFTTRKLPKVLEVDRNTYIKRIILDSEENLLGCESLNFMRSYSDNNLQT- 181" FVPESFDRDKTIALIMNSGSTGLPKGVALPHRTACVRFSHARDPIFGNQIPTDIALS **., **, **, **, **, **, **, **, **, **, **, ** 177" FKPLHYDOPDVQAAILCSSLGTTLPKGVMQTHQNICVRILHALDPRTQQLIPGVSVL 241" VPFFHGFGMFTLGYLICGRVRLMYRFEELFLRSLODQYIQSALLVPTLSSFAKSTL ***, **, **, **, **, **, **, **, **, **, **, ** 237" LPFFHAFGSINLGYFMVGLRVMFRRFDQEAFLKAIDQYEVRSINVPSTILFLSKSP 301" IDKYLDSLNLHEIASGGAPLSKEVEAVALKRFLPGLPQYQGLTETTSAILITPEGDDKP *****+, **, **, **, **, **, **, **, **, **, **, ** 297" VDKYDLSTLAECLCGAAPLAKEVAEAAKRNLPGIRCGFLTESTSAIHTLHNEFKSG 361" AVGVVPFFEAKWVLDLTGKTLGVNQRGECLVRGPIMISGYVNNPEATNALIDDGWLH ***, **, **, **, **, **, **, **, **, **, ** 357" SLGKVTYMAAKIADRNTEGALGPQNQVGECLIKGPMVTKYVNNPQATKEAIDDDGWLS 421" GDIAYWDEDEHFFIVDRLKSLIKYKGYQVAPAELESILQHPNIFDAGVAGLPDDDAEGE ***, **, **, **, **, **, **, **, **, **, ** 417" GDGFYDDEDEHFYVDRYKELIKYKGQVAPVELEILLQHPGIRDVAVGIPDIEAGEL 481" PAAVVLLEHGTMTKEIVDVAQSQTAAKKLRGGV/FVDEVPKGLTGKLDARK-IREI **., **, **, **, **, **, **, **, **, **, ** 477" PAGFVVKQPGTEITAKEVDYLAERVSHTKYLRRGVRFVDSIPRNVTGKTRKELLKQL 541" KAKGGKIAV 537" EKASKL	[81.3% / 540 aa] 1" MERENKNVYGPPEPKHPLGNFTAGEMLYNALHKHSHHQI-LDVMGNESLSYQEFFDTT ...*, **, **, **, ****, ****, *, ..., *., *, ...*, ** 1" MVKREKNVIYGPPEPLHPLEDLTAGEMLFRALRKHSHL-P-AOLDVVGDESLYKEFFEA 60" KLGSQSLQNCGYKMNDVWSICAENNRKFIPISAWSYIGMIVAPVNEDYIPDELCKVTG ...*, **, **, **, **, **, **, **, **, **, ** 61" LLAQSLHNCGYKMNDVWSICAENNRKFIPVAAWYIGMIVAPVNEDYIPDELCKVMG 120" KPLIPVFTTRKLPKVLEVDRNTYIKRIILDSEENLLGCESLNFMRSYSDNNLQTFKP **., **, **, **, **, **, **, **, **, **, ** 121" QKPVQIVFTTKNILNKVLEVOVSRTNFIRIILDTVENIHGCESLPNF-SRYSDGNIANFKP 180" LHYPDVQVAAILCSSLGTTLPKGVMQTHQNICVRILHALDPRTQQLIPGVSVL ...*, **, **, **, **, **, **, **, **, **, ** 181" LHPDVPVQEVAAILCSSLGTTLPKGVMQTHQNICVRILHALDPRTQQLIPGVTVL 240" FAHAFGSINLGYFMVGLRVMFRRFDQEAFLKAIDQYEVRSINVPSTILFLSKSP *****+, **, **, **, **, **, **, **, **, **, ** 241" FAHAFGHITLGYFMVGLRVMFRRFDQEAFLKAIDQYEVRSINVPSTILFLSKSP 300" YDLSTLAECLCGAAPLAKEVAEAAKRNLPGIRCGFLTESTSAIHTLHNEFKSG *****+, **, **, **, **, **, **, **, **, **, ** 301" YDLSSRELCCGAAPLAKEVAEAAKRNLPGIRCGFLTESTSAIHTLHNEFKSG 360" KVTPYMAAKIADRNTEGALGPQNQVGECLIKGPMVTKYVNNPQATKEAIDDDGWLS ***, **, **, **, **, **, **, **, **, **, ** 361" RVTPLMAAKIADRETGKALGPQNQVGECLIKGPMVSKYVNNVTEAKIAIDDDGWLS 420" GYYDEDEFYIVDRYKELIKYKGYQVAPVELEILLQHPGIRDVAVGIPDIEAGELPAG *****+, **, **, **, **, **, **, **, **, **, ** 421" GYYDEDEFYIVDRYKELIKYKGQVAPAELEEILKNPCIRDVAVGIPDLEAGELPSA 480" FVVKQPGTEITAKEVDYLAERVSHTKYLRRGVRFVDSIPRNVTGKTRKELLKQL ...*, **, **, **, **, **, **, **, **, **, ** 481" FVVKQPGTEITAKEVDYLAERVSHTKYLRRGVRFVDSIPRNVTGKTRKELLKQL 540" SKL .GG 541" GG