
FGSC# my sg No. Origin: Mutagenesis or segregants (or recombinants) from diploids

- A967 = EK157 from 2n (**025**) : EK106 = N907 = FGSC#A942 / EK113 from 2n (**002**)
EK113 = cspA1; (acrA1)* brnA2; choA101; nicA1; pdxA2; trpB2
 2n (**002**) : N837 = FGSC#A925 = EK040 / EK053 = FGSC# A954 (above)
- A968 = EK185 from 2n (**042**) : (see FGSC#A960, p.1)
- A969 = EK182 from 2n (**041**) : N814 = FGSC#A920 = [EK034] / N639 = FGSC#A897 = EK037
- A970 = EK187 from 2n (**042**) : (see FGSC#A960, p.1)
- A971 = EK184 from 2n (**041**) : (see FGSC#A969, above)
- A972 = EK191 from 2n (**043**) : N814 = FGSC#A920 / EK135 [from 2n (**008**), see A964, p.1]
- A973 = EK199 from 2n (**037**) : EK171 (see A961, p.1) / EK164 = FGSC#A957 (see p.1)
- A974 = EK210 X-ray of EK146 --> *sftC101*
EK146 = cspA1; fwnA1; leuA1; pdxA2, from 2n (**016**) [see FGSC#A958, p.1]
- A975 = EK219 from 2n (**052**) : EK210 = FGSC#A974 / EK175 (from 2n (**031**))
EK175 = cspA1; (acrA1)* brnA2; fpaD19; choA101; metB11; oliC2; crnB12
 2n (**031**) : EK138 = FGSC#A978 / N784 = EK131
 N784 = cspA1; fwnA1; **fpaD19**; lysA14 (Debets. unpubl.; see FGSC#A961, p. 1)
- A978 = EK138 from 2n (**023**) : EK106 = N907 = FGSC#A942 / EK112 from diploid (**002**)
EK112 = cspA1; (acrA1)* brnA2; choA101; nicA1; pdxA2; trpB2
 2n (**002**) : EK040 = N837 = FGSC#A925 / EK053 = FGSC#A954 (see p. 1)
- A979 = EK192 from 2n (**032**) : EK172 / EK171 ; both, 172 & 171 from diploid (**031**); see A961, p.1]
EK171 = cspA1; (acrA1)* brnA2; fpaD19; lysA14; metB11; oliC2,
EK172 = cspA1; fwnA2; argH12; choA101; metB11; pdxA2; crnB12
- A980 = EK194 from 2n (**032**) [see preceding A979, and also FGSC#A975, above]
- A981 = EK201 from 2n (**036**) : EK171 / EK158 = FGSC#A956, p. 1)
EK171 from diploid (**031**) [see preceding FGSC#A979 & A961]
- A982 = EK223 }
A983 = EK224 } all from 2n (**051**) : EK200 = FGSC#A962 (p.1) / EK193 from diploid (**032**)
A984 = EK225 }
A985 = EK226 }
EK193 = cspA1; (acrA1)* brnA2; fpaD19; choA101; metB11; pdxA2; oliC2; crnB12
 2n (**032**) : EK172 / EK171 [see above, FGSC#A979]

* (acrA1) indicates the likely presence of acrA1, closely linked to brnA2, but not confirmed by growth tests