Cockatiel mutations

Courtesy of The Sunraysia Branch of The Aviculture Sosiety of Australia **Remember when it was a dark grey bird, below is a list of mutations as identified by the Australian National Cockatiel Society and used as a SHOW STANDARD.**

Normal

Body to be dark grey, uniform throughout. Males to have a bright clear head of deep yellow, with bright orange cheek patches. Wing patch to be well defined and clean of darker feathers. Females to be of duller version of male colouration with barring on tail which is absent in males.

Pearl

Hens. Well defined heavy pearling is the aim with good depth of colour in pearls and body. Cocks. As for hens with less importance placed on pearl quality.

Lutino

Ideally a uniform rich deep lemon yellow colour throughout with bright contrasting cheek patches of orange. Flesh coloured feet and clear nails. Eyes to be ruby or red.

Cinnamon

Rich uniform cinnamon colouration on entire body, with yellow head area and bright cheek patch. No blotching or mottled effect. Flesh coloured feet, clear nails and dark ruby eyes.

Heavy Pearl-Pied

Same as for heavy pied with even pearl feather pattern on the non pied markings Light Pearl-Pied

Same as for light pied with even pearl feather pattern on the non pied markings

Fallow

A light cinnamon colour suffused by yellow, which is very evident on chest. The eyes to be clear red. Flesh coloured feet and nails.

Silver(Recessive)

A diluted form of the normal body colour which is best described as silver grey. Colour to be uniform throughout. Eye colour is lighter than normal

Whiteface

As per normal but totally devoid of any yellow or orange colouration. Birds to be two colours only dark grey and white. Cocks to have pure white head with no dark feathers and good definition between head and body colour.

Platinum

A pale smoky grey colour with a yellow suffusion throughout body. Yellow head area and bright cheek patch. The smoky grey colour is slightly darker on tail and flights. Feet and nails flesh coloured.

Multi mutations

Are many and varied and will be judged by combining the mutation specifications relevant to the particular multi mutation