How to insert or replace a bird in DMS – *even on a saved card*

Great news for users of DMS

If you are compiling a card and suddenly realise that you have missed an entry way down near the beginning of your list *don't despair!* I had thought that the only way to fix this was to delete all the entries above and start again from where you make the error, a tedious business if the Whyte's Barbet should have been number 2 and you are now at number 86…! And here I should admit that in the past I have omitted a bird and thought it not worth the effort to redo it all and have simply sent it in 'as is'.

But here's how to do it. In the example I have just done a short card to illustrate the method and need to **replace** the Sombre Greenbul (No. 1) with Dark-capped Bulbul. This is a card that hasn't been saved.

Search for bulbuls to get the list opposite.

Select Dark-capped, change *Surveyed Sequence* to 1 in this case (it would have shown 5 as the next in the list), and click on *Add*.

Sequence	Spp	SppName
4	552	Kurrichane Thrush
3	580	White-browed Robin-Chat*
2	337	Purple-crested Turaco*
1	545	Dark-capped Bulbul*
1	551	Sombre Greenbul

Sequence	Spp	SppName	
4	552	Kurrichane Thrush	
3	580	White-browed Robin-Chat*	
2	337	Purple-crested Turaco*	
1	545	Dark-capped Bulbul*	

You now have two 1's (left). Just click on the sombre or even the *Sequence* number 1 and click *Delete species*. The bird is replaced as shown on the right!

To **insert** a missed bird the process is similar. Here I insert Bar-throated Apalis as number 3.

Sequence	Spp	SppName
4	552	Kurrichane Thrush
3	622	Bar-throated Apalis*
3	580	White-browed Robin-Chat*
2	337	Purple-crested Turaco*
1	545	Dark-capped Bulbul*

Select the apalis and enter the *sequence* number as number 3.

Click *Add*.

Your new sequence will bring up an error when you **save the card**.

Sequence	CardNo	Spp	SppName	ID	Sent
4	1745_3100a005476a20130904	552	Kurrichane Thrush	10024	0.0000
3					
3					
2					
1					

Errors have been found on this form. Please correct them before trying to save t... ✕

You have a duplicate in the species sequence at sequence number: 3

OK

☐ **This card covers a minimum initial** [Add Additional Info] [Correct sequence] [Delete species]

A new box has appeared so click OK and then *Correct sequence*.

Sequence	CardNo	Spp	SppName
5	1745_3100a005476a20130904	552	Kurrichane Thrush
4	1745_3100a005476a20130904	580	White-browed Robin-Chat*
3	1745_3100a005476a20130904	622	Bar-throated Apalis*
2	1745_3100a005476a20130904	337	Purple-crested Turaco*
1	1745_3100a005476a20130904	545	Dark-capped Bulbul*

The card is saved and the apalis added as no. 3 with the correct sequence.

No problems!

You might get an ORF because you haven't picked up that you have entered a wrong name. This is a fairly frequent error that is corrected by notifying raczim and done on the Sabap2 website – if you are using the on-line system you will have to stick to this method as I don't think you can correct your own card.

On DMS you can do the correction yourself. Browse your records to the relevant card; I'll use the one above as an example. The layout of a saved card looks like this…

CardNo	Spp	SppName	Sequence	Sent	ID
1745_3100a005476a20130904	545	Dark-capped Bulbul*	1	0.0000	9306
1745_3100a005476a20130904	337	Purple-crested Turaco*	2	0.0000	9307
1745_3100a005476a20130904	622	Bar-throated Apalis*	3	0.0000	9310
1745_3100a005476a20130904	580	White-browed Robin-Chat*	4	0.0000	9308
1745_3100a005476a20130904	552	Kurrichane Thrush	5	0.0000	9309

Let's say the Bar-throated Apalis is an ORF and should have been a Yellow-breasted. The *Surveyed sequence* box will be blank so select Yellow-breasted, enter 3 in the box and *Add*. Then just click on Bar-throated and *Delete species*. The card is changed – save it and re-submit it. You could also insert a bird you previously missed, following the above procedure. Make sure the **Totals** box reflects the changes. You will get another error message is it hasn't kept up with your changes.

Thanks to Celesta, via Geoff Lockwood, for bringing this useful method to my attention.

Happy atlassing

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