

CCPMT Step 2 - Microsoft Internet Explorer

File Edit View Favorites Tools Help

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Address http://localhost/ccpmt/Ccpmt-Step2-new.jsp?ID=probesetId&input_array=Select&email=ruchi@uab.edu&B1=Go+to+Step+2 Go Links

Google Search 1698 blocked ABC Check AutoLink AutoFill Options

Step 1

Select the input for your mapping.

Step 2

Select the output you want mapped

OUTPUT - Step 2

- Enter the IDs for mapping in the box

244904_at

(e.g. for AGI: AT3G26650,AT1G09970)
 (e.g. for Arabidopsis Probesets: 14686_at,13080_at)
 (e.g. for EGO: 893982,915242)
 (e.g. for TCs: TC251326,TC31967)
- Select the species type for the input values

Arabidopsis
- Select the output arrays to be mapped

Arabidopsis	Poplar
<input checked="" type="checkbox"/> Affymetrix Arabidopsis Genome (8k) <input checked="" type="checkbox"/> Affymetrix Arabidopsis Genome ATH1(25K) <input type="checkbox"/> AFGC Arabidopsis Array <input type="checkbox"/> Operon Arabidopsis Genome Oligo <input type="checkbox"/> Agilent Arabidopsis 2 Oligo <input type="checkbox"/> CATMA - Complete Arabidopsis Transcriptome	<input type="checkbox"/> Affymetrix Poplar Genome Array

Done Local intranet