## Attribute Mapping within RippleDown

RippleDown will automatically add any new attributes received by the data source to the relevant project. The attribute mapping for all projects can be found in RippleDown by:

1. Logging in to the Knowledge Builder module.

<b><i><b>Ripple</b></i></b>	Dov	wn
Version: 8.0.5		
Caboratory Systems Calidator Calidator Calidator Calidator		
Business Audit Systems O Data Entry Auditor		
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Name Password		]
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- 2. Open the desired project.
- 3. Go to Edit | Attributes.

쭏 Knowledge Builder - RippleDown



4. The attribute editor will display all attributes ever received by RippleDown for the installation instance. The display name within RippleDown will be shown.

Attribute Editor       X         File       Edit       View       Options         File       Edit       View       Options         Very Type here to filter the tree       Image: Comparison of the tree       Image: Comparison of the tree         HDL_chol       Triglycerides       Image: Comparison of the tree         HDL_chol       Tmage: Comparison of the tree       Image: Comparison of the tree         HDL_chol       Tmage: Comparison of the tree       Image: Comparison of the tree         HDL_chol       Tmage: C		
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Q- Type here to filter the tree         HDL_chol         LDL_chol         Triglycerides         Total Chol HDL Ratio         LDL_HDL Ratio         LDL_HDL Ratio         LDL_HDL Ratio         APO A1         APO B         APO A1         APO B         APO C GENOTYPE         BROAD BETA         LDL_FPG         VLDL_EPG         ALP         AST         GGT         Albumin         Total Protein		
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VLDL_EPG ALP ALP_Marked_Elev ALT AST GGT Albumin Total Protein	LDL EPG	
ALP ALP_Marked_Elev ALT AST GGT Albumin Total Protein Y Close	VLDL EPG	
ALP_Marked_Elev ALT AST GGT Albumin Total Protein      Close     @ Help	ALP	
ALT AST GGT Albumin Total Protein	ALP Marked Elev	
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Close O <u>H</u> elp	Total Protein	
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This document is intended to be used as a supplement to the RippleDown Administrator Guide and RippleDown Knowledge Builder Guide. http://support.pks.com.au/product-register/ 5. From the Options menu select "Show external names".



6. The external name mapped to each attribute will now be shown beside the display name.

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<u>F</u> ile <u>E</u> dit <u>V</u> iew <u>O</u> ptions	
Q- Type here to filter the tree	
HDL_chol (Hd)	-
LDL_chol (LD)	
Triglycerides (Tr)	
Total Chol_HDL Ratio (Us)	
LDL_HDL Ratio (LH)	
APO A1 (Z7)	
APO B (Z8)	
Apo B_APO A1 RATIO (Z9)	
Lipoprotein_a (Oo)	
APO E GENOTYPE (Y3)	
BROAD BETA (Y5)	
LDL_EPG (Y7)	
VLDL_EPG (Y6)	
ALP (Ap)	
ALP_Marked_Elev	
ALT (At)	
AST (As)	
GGT (Gg)	
Albumin (Ab)	
Total Protein (Tp)	¥
Close @ H	elp

7. The external name of an attribute can be changed first selecting the attribute, then selecting "Set external name" from the Option menu.

河 Attribute Editor		×
<u>File Edit View</u>	Options	
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O True have to fi	S <u>e</u> t external name	
HDL chol (Hd)	Highlight colour	
LDL_chol (LD)	<u>S</u> et default format	•
Triglycerides (Tr	)	<i></i>
Total Chol HDL	Ratio (Us)	

Set external name	for "HDL_chol"		×
External name	Hd		
		<u>o</u> ĸ	Cancel

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