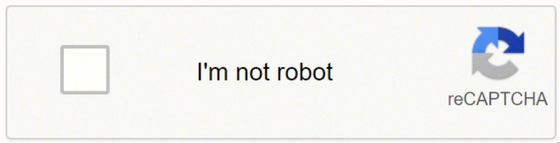

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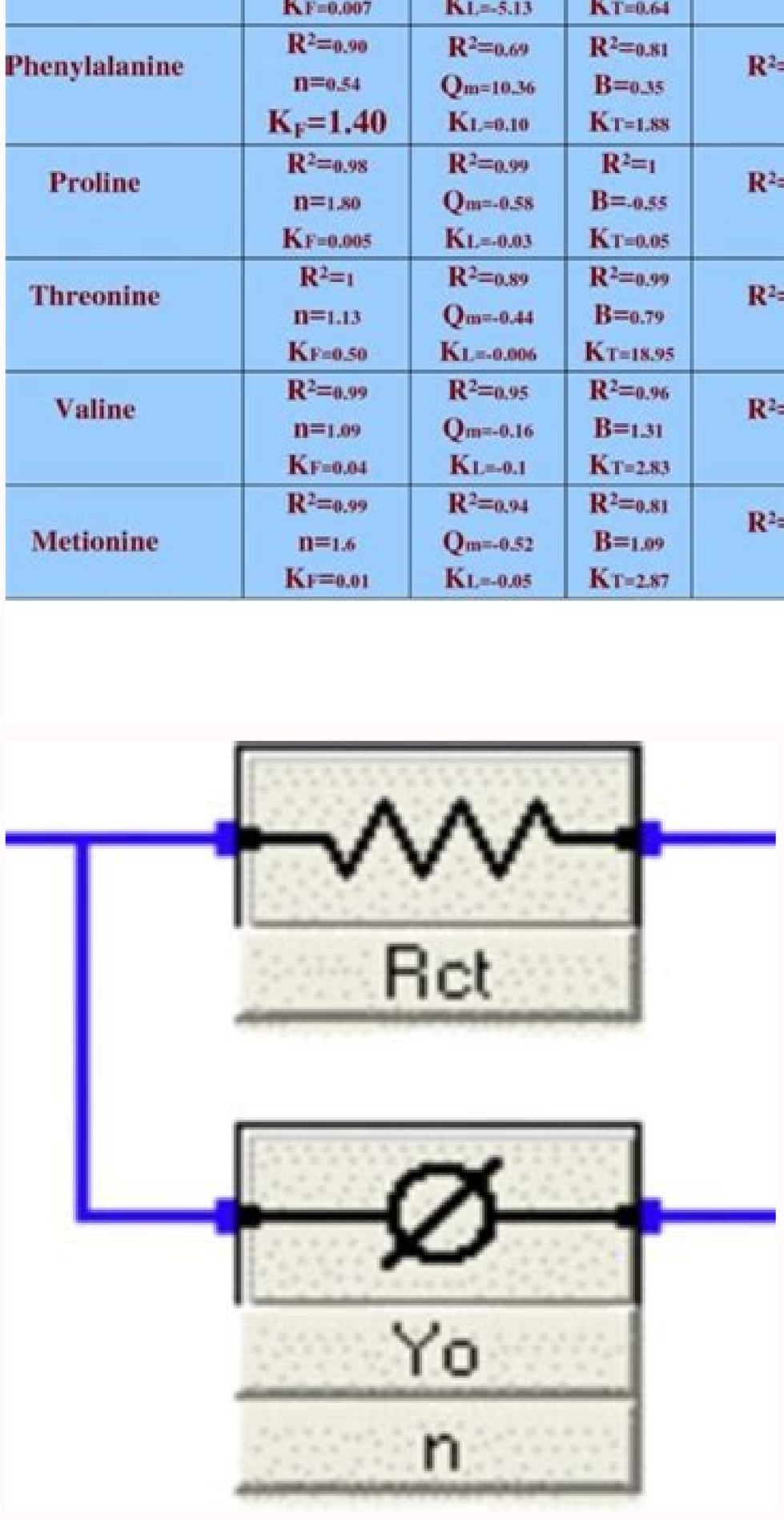
28570568110 14436723.388889 6138819.1076923 34197176740 751920651 83859474744 21750713578 140278822 19290995.463768 6702868998 95403064680

sotherm parameters	Parameter value
.angmuir	
$_m$(mg g^{-1})	115
$_L$(L mg^{-1})	1.2x10^{-3}
^{2}	0.978
reundlich	
	2.70
$_F$(mg g^{-1})(L mg^{-1})$^{1/n}$	10.22
^{2}	0.969
emkim	
(L mg^{-1})	0.307
	18.27
$_T$(J mol^{-1})	137
^{2}	0.934
Dubinin-Redushkevich	
$_{DR}$(mg g^{-1})	178.7
(Mol J^{-1})^2	4.24x10^{-9}
(kJ Mol^{-1})	10.85
^{2}	0.976

Cystein	**n=0.96** **K_F=0.58**	**Q_m=22.37** **K_L=0.03**	**B=0.26** **K_T=1.69**	**R^2=0.44**
Glycine	**R^2= 0.99** **n=1.88** **K_F=0.10**	**R^2=0.90** **Q_m=-0.78** **K_L=-0.14**	**R^2=0.98** **B=0.60** **K_T=1.55**	**R^2=0.94**
Histidin	**R^2=0.96** **n=1.5** **K_F=0.007**	**R^2=0.61** **Q_m=2.07** **K_L=-5.13**	**R^2=0.99** **B=0.73** **K_T=0.64**	**R^2=0.87**
Phenylalanine	**R^2=0.90** **n=0.54** **K_F=1.40**	**R^2=0.69** **Q_m=10.36** **K_L=0.10**	**R^2=0.81** **B=0.35** **K_T=1.88**	**R^2=0.82**
Proline	**R^2=0.98** **n=1.80** **K_F=0.005**	**R^2=0.99** **Q_m=-0.58** **K_L=-0.03**	**R^2=1** **B=-0.55** **K_T=0.05**	**R^2=0.94**
Threonine	**R^2=1** **n=1.13** **K_F=0.50**	**R^2=0.89** **Q_m=-0.44** **K_L=-0.006**	**R^2=0.99** **B=0.79** **K_T=18.95**	**R^2=0.87**
Valine	**R^2=0.99** **n=1.09** **K_F=0.04**	**R^2=0.95** **Q_m=-0.16** **K_L=-0.1**	**R^2=0.96** **B=1.31** **K_T=2.83**	**R^2=0.89**
Metionine	**R^2=0.99** **n=1.6** **K_F=0.01**	**R^2=0.94** **Q_m=-0.52** **K_L=-0.05**	**R^2=0.81** **B=1.09** **K_T=2.87**	**R^2=0.90**



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[PubMed] [CrossRef] [Google Scholar]Page 2Identified functional groups present in Saccharum arundinaceum by FTIR spectroscopy.Possible functional groupsRaw (cm−1)Base treated (cm−1)Acid treated (cm−1)–OH stretching3309–37513211–3400 =CH 3294C–H2858–2918 (bifurcate)29102922C=O Secondary amide164516081654.92–NH C–O11071222 C–N105110561043 Malandrino M., Abollino O., Giacomino A., Aceto M., Mentasti E. doi: 10.1016/j.seppur.2007.10.002. 1990;24(1):112–118. 2001;35(5):1125–1134. D., Spiridon O. 2015;2(1):36–46. 2013;8(1) doi: 10.15376/biores.8.1.1147-1165. Iqbal M., Saeed A., Zafar S. FTIR spectrophotometry, kinetics and adsorption isotherms modeling, ion exchange, and EDX analysis for understanding the mechanism of Cd2+ and Pb2+ removal by mango peel waste. doi: 10.1007/s11783-009-0030-7. doi: 10.1016/s0043-1354(00)00389-4. Colloids and Surfaces B: Biointerfaces. doi: 10.1016/j.watres.2005.10.040. doi: 10.1016/j.jcis.2006.03.011. doi: 10.1016/j.desal.2005.10.032. 2008;144(2):235–244. 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