2013 Joint Conference of Drug Safety Research Centres In affiliation with the Pacific Rim Association for Clinical Pharmacogenetics (PRACP)

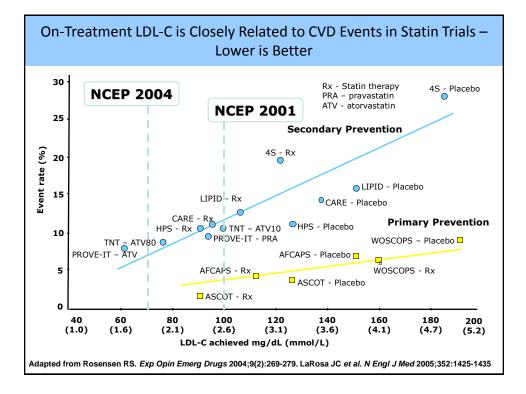
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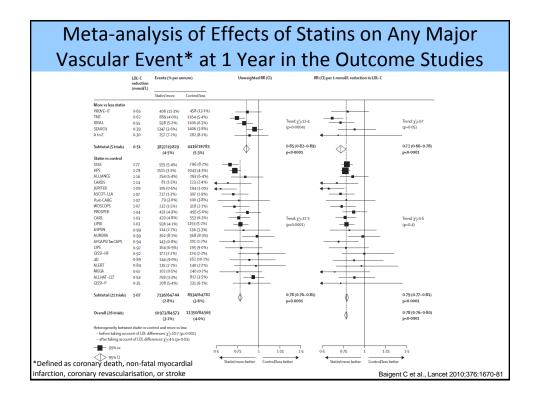
Optimising the Efficacy & Safety of Statins

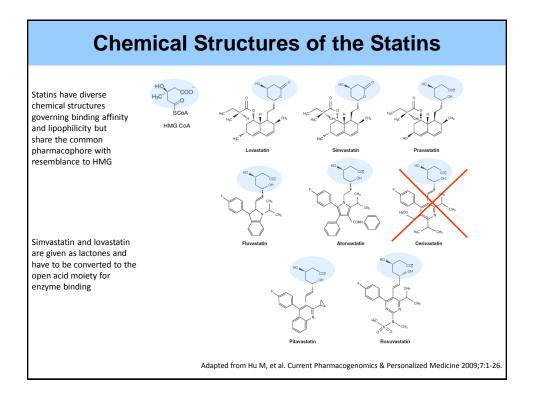


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Statins and risk of incident diabetes: a collaborative meta-analysis of randomised statin trials

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Naveed Sattar, David Preiss, Heather M Murray, Paul Welsh, Brendan M Buckley, Anton J M de Craen, Sreenivasa Rao Kondapally Seshasai, John J MA/lurray, Dilys J Freeman, J Wouter Jukema, Peter W Madfarlane, Chris J Packard, David J Stott, Rauf G Westendorp, James Shepherd, Barry R Davis, Sara L Pressad, Roberto Marchioli, Rosa Maria Marfisi, Aldo P Maggioni, Luigi Tavazzi, Gianni Tognoni, John Kjekshus, Terje R Pedersen, Thomas J Cook, Antonio M Gotto, Michael B Clearfield, John R Downs, Haruo Nakamurru, Yasuo Ohashi, Kyoichin Mizuno, Nausik K Ray, Ian Ford

Summary

Background Trials of statin therapy have had conflicting findings on the risk of development of diabetes mellitus in patients given statins. We aimed to establish by a meta-analysis of published and unpublished data whether any relation exists between statin use and development of diabetes.

Methods We searched Medline, Embase, and the Cochrane Central Register of Controlled Trials from 1994 to 2009, for randomised controlled endpoint trials of statins. We included only trials with more than 1000 patients, with identical follow-up in both groups and duration of more than 1 year. We excluded trials of patients with organ transplants or who needed haemodialysis. We used the *P* statistic to measure heterogeneity between trials and calculated risk estimates for incident diabetes with random-effect meta-analysis.

Findings We identified 13 statin trials with 91140 participants, of whom 4278 (2226 assigned statins and 2052 assigned control treatment) developed diabetes during a mean of 4 years. Statin therapy was associated with a 9% increased risk for incident diabetes (odds ratio [OR] 1-09; 95% CI 1-02–1-17), with little heterogeneity (*I*=11%) between trials. Meta-regression showed that risk of development of diabetes with statins was highest in trials with older participants, but neither baseline body-mass index nor change in LDL-cholesterol concentrations accounted for residual variation in risk. Treatment of 255 (95% CI 150–852) patients with statins for 4 years resulted in one extra case of diabetes.

Interpretation Statin therapy is associated with a slightly increased risk of development of diabetes, but the risk is low both in absolute terms and when compared with the reduction in coronary events. Clinical practice in patients with moderate or high cardiovascular risk or existing cardiovascular disease should not change.

Published Online February 17, 2010 DOI:10.1016/S0140-6726(09)61965-6 See Comment page 700 British Heart Foondation Glasgow Cardiovascular Research Centre, University of Glasgow, UK (Porl N Sattar PhD, D Preiss MRCP, PWelsh PhD, Prof.] J McMurray MD); Robertson

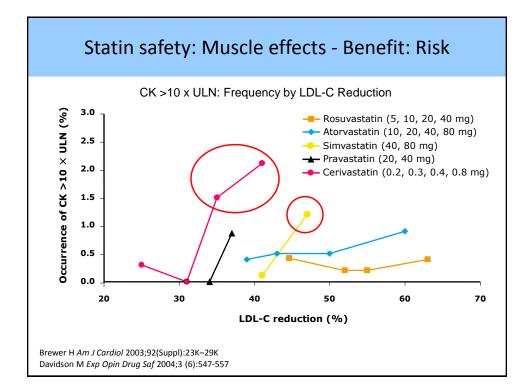
Prof J McMurray MUJ; Robertson Centre for Blockstatikis, University of Glasgow, Glasgow, UK (HIM Murray MSc, Prof I Ford PhD); Department of Pharmacology and Therapeutics, Cork, University Hospital, Cork, Ireland (Hof B Macdaky FRCH); Department of Gerontology and Genatrics, Laledu University Medical Contre, Leiden, Netherlands

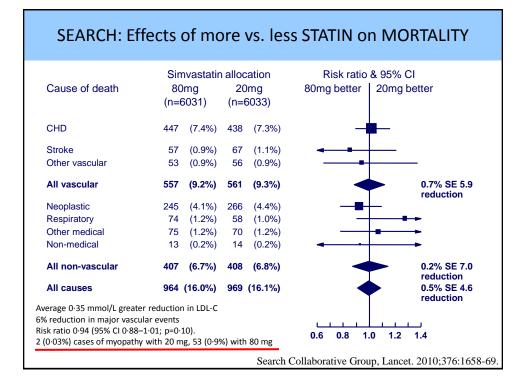
Stattar N, et al., Lancet 2010;375:735-42.

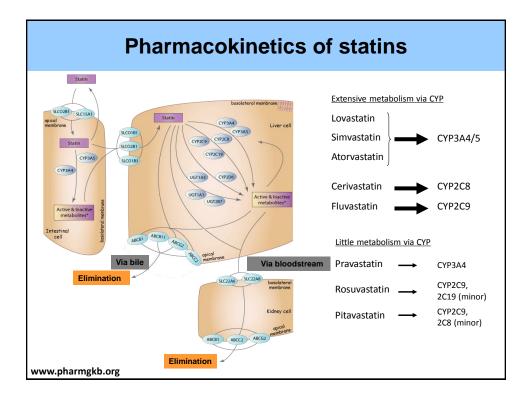
Muscle toxicity

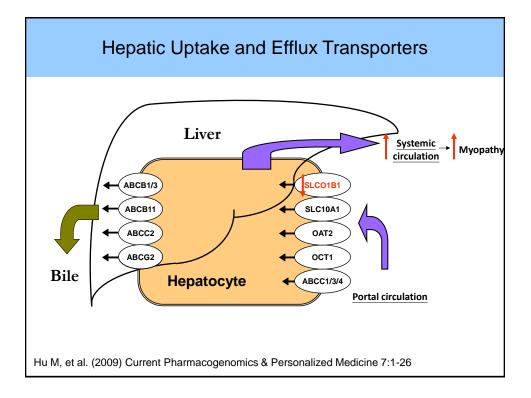
- Muscle toxicity is the most significant and well documented adverse effect with statins
- Mild symptoms of muscle pain, tenderness and weakness are relatively common, but fortunately, the most severe and potentially fatal condition of rhabdomyolysis is very rare
- All statins can cause muscle toxicity if given in high enough dosage or if there are drug interactions increasing systemic exposure
- Several predisposing factors have been identified, including old age, renal/liver dysfunction and hypothyroidism, and some patients have a genetic predisposition which either alters the systemic exposure to the drug or causes susceptibility in the muscles

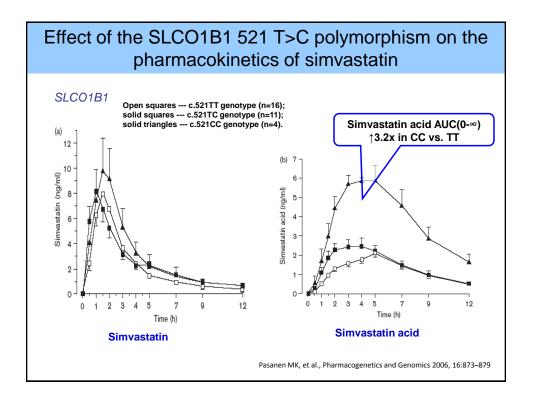
Hu M, Cheung BMY, Tomlinson B. Ther Adv Drug Saf 2012;3(3): 133-144

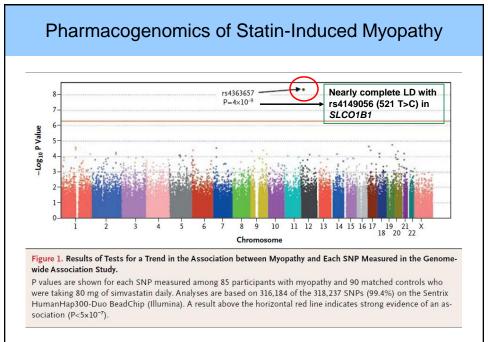




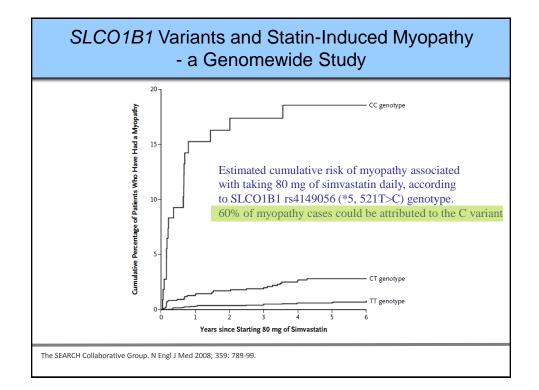








The SEARCH Collaborative Group. N Engl J Med 2008; 359: 789-99.



FDA guideline on 8 June 2011 on limiting the use of simvastatin 80 mg

FDA U.S. Food and Drug Administration

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News & Events

FDA NEWS RELEASE

For Immediate Release: June 8, 2011 Media Inquiries: Morgan Liscinsky, 301-796-0397, morgan.liscinsky@fda.hhs.gov

Consumer Inquiries: 888-INFO-FDA FDA announces new safety recommendations for high-dose simvastatin

Increased risk of muscle Injury cited

The U.S. Food and Drug Administration today is announcing safety label changes for the cholesterol-lowering medication simvastatin because the highest approved dose--80 milligram (mg)--has been associated with an elevated risk of muscle injury or myopathy, particularly during the first 12 months of use.

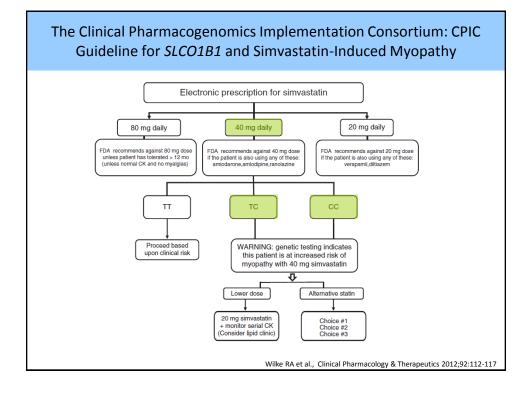
The agency is recommending that simvastatin 60 mg be used only in patients who have been taking this dose for 12 months or more and have not experienced any muscle toxicity. It should not be prescribed to new patients. There are also new contraindications and dose limitations for when simvastatin is taken with certain other medications.

Simvastatin is used together with diet and exercise to reduce the amount of "bad cholesterol" (low-density lipoprotein cholesterol or LDL-C) in the blood. High levels c LDL-C are linked to a higher risk of heart attack, stroke and cardiovascular death. In 2010, about 2.1 million patients in the United States were prescribed a product containing simvastatin 80 mg.

"The FDA has completed its review of the safety of high-dose sinvastatin and is making label changes to reduce the risk of statin-associated muscle injury," said Eric Colman, M.D., deputy director of the Division of Metabolism and Endocrinology Products in the FDA's Center for Drug Evaluation and Research. "We want to ensure that patients and health care professionals are aware of the new labeling changes to simvastatin, including the increased risk of myopathy when using the 80 mg dost of simvastatin."

The changes to the label for simvastatin-containing medications are based on the FDA's review of the results of the seven-year Study of the Effectiveness of Additiona Reductions in Cholesterol and Homocysteine clinical trial, other clinical trial data, and analyses of diverse events submitted to the FDA's Adverse Event Reporting System. All showed that patients taking aliversatin 80 mg daily had an Increased risk of muscle Injury compared to patients taking lower does of simusatatin or other statin drugs. The risk of muscle Injury is highest during the first year of treatment with the 80 mg doe of simvastatin, is often the result of Interactions with certain other medicines, and is frequently associated with a genetic predictiposition for simvastatin-related muscle Injury.

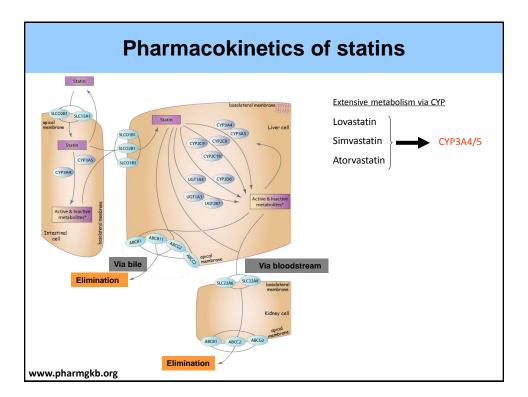
Simvastatin is sold under the brand-name Zocor and as a single-ingredient generic product. It is also sold in combination with ezetimibe as Vytorin and in combination with niacin as Simcor.



Effect of SLCO1B1 521T>C polymorphism on the lipid-lowering effect of statins

- Each copy of the 521C variant allele was only associated with -1.28% smaller reductions in LDL-C in response to simvastatin in the Heart Protection Study (>10,000 patients)
- No significantly noticeable effect on the lipid-lowering effect of statins

The SEARCH Collaborative Group. N Engl J Med 2008; 359: 789-99. Hu M, Tomlinson B. *Pharmacogenomics*. 2013 Jun;14(8):981-95.



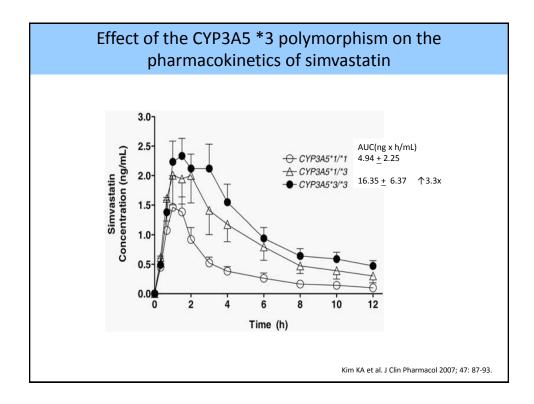
Clinical Pharmacology & Therapeutics

PPARA: A Novel Genetic Determinant of CYP3A4 *In Vitro* and *In Vivo*

Kathrin Klein¹, Maria Thomas¹, Stefan Winter¹, Andreas K. Nussler², Mikko Niemi³, Matthias Schwab^{1,4} and Ulrich M. Zanger¹

Interindividual variability in cytochrome P450 3A4 (CYP3A4) is believed to be largely heritable; however, predictive genetic factors have remained scarce. Using a candidate-gene approach in a human liver bank, we identified singlenucleotide polymorphisms (SNPs) in the Ah-receptor nuclear translocator (*ARNT*), glucocorticoid receptor (GR), progesterone receptor membrane component 2 (*PGRMC2*), and peroxisome proliferator-activated receptor-a (*PPARA*) that are associated with CYP3A4 phenotype. Validation in atorvastatin-treated volunteers confirmed a decrease in atorvastatin-2-hydroxylation in carriers of *PPARA* SNP rs4253728. Homozygous carriers expressed significantly less PPAR-a protein in the liver. Moreover, shRNA-mediated *PPARA* gene knockdown in primary human hepatocytes decreased expression levels of the PPAR-a target ACOX1 and of CYP3A4 by more than 50%. In conclusion, this study identified novel genetic determinants of CYP3A4 that, together with nongenetic factors, explained 52, 55, and 33% of hepatic CYP3A4 mRNA, protein, and atorvastatin-2-hydroxylase activity, respectively. These findings have implications for variability in response to drug substrates of CYP3A4.

Clinical Pharmacology & Therapeutics (2012); 91 6, 1044–1052





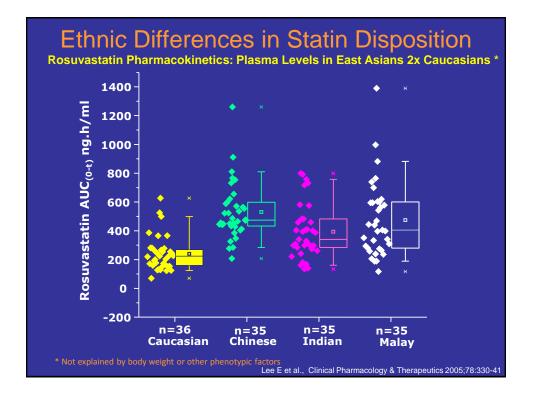
Associations between the genotypes and phenotype of CYP3A and the lipid response to simvastatin in Chinese patients with hypercholesterolemia

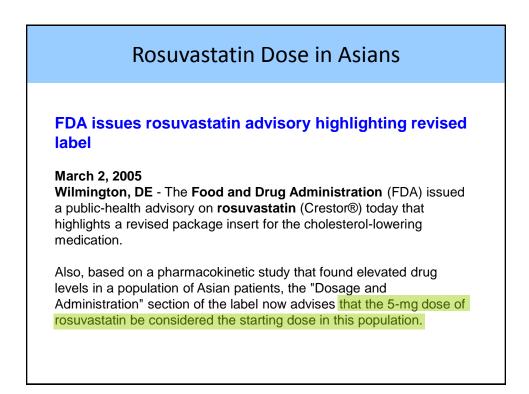
Aim: This study examined the associations between the CYP3A4*1G, CYP3A4*22, CYP3A5*3 and PPARA rs4823613A>G polymorphisms and the phenotypes of CYP3A estimated by the ratio of 6 β -hydroxycortisol:cortisol in urine, and the low-density lipoprotein cholesterol response to simvastatin in Chinese patients with hypercholesterolemia. Patients & methods: Lipid profiles were determined off treatment and after 6 weeks of treatment with simvastatin 40 mg in 273 patients. Results: There was no significant association between the ratio of 6 β -hydroxycortisol:cortisol and the low-density lipoprotein cholesterol response to simvastatin in the study subjects (r = 0.052; p = 0.455). The genetic polymorphisms examined had no significant association with this measure of CYP3A phenotype or the lipid-lowering responses to simvastatin. Conclusion: The results of this study suggest that genetic polymorphisms in CYP3A or other regulatory genes, or the CYP3A activity itself, is unlikely to have a significant effect on the lipid-lowering responses to simvastatin in Chinese patients.

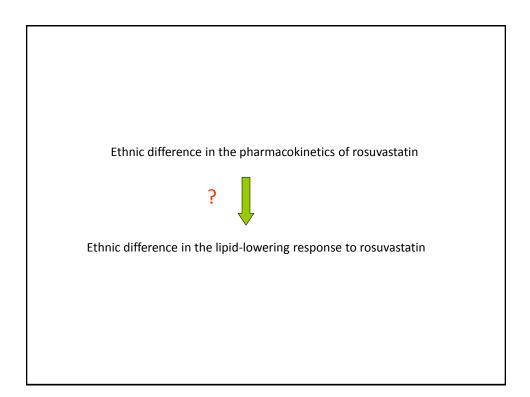
Hu M,.....Tomlinson B. Pharmacogenomics (2013) 14(1), 25-34

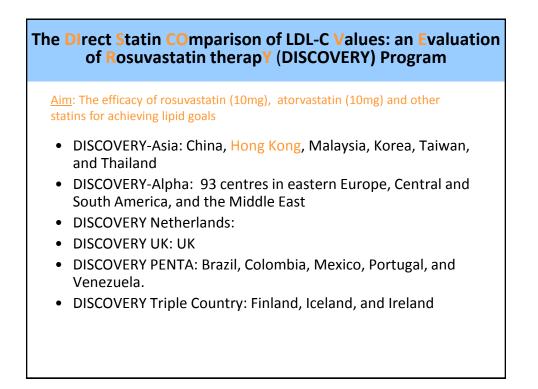
Genetic polymorphisms in CYP3A or PPARA had no significant effect on the lipid-lowering responses to simvastatin

Genetic polymorphisms	Genotypes	LDL-C response to simvastatin (%)	
CYP3A4*1G	*1*1 (n = 143)	-47.5 ± 11.6	
	*1*1G (n = 107)	-46.7 ± 10.9	<i>P</i> >0.05
	*1G*1G (n = 18)	-47.7 ± 14.8	
СҮРЗА5*3	*1*1 (n = 18)	-50.2 ± 9.6	
	*1*3 (n = 107)	-47.2 ± 12.1	<i>P</i> >0.05
	*3*3 (n = 144)	-47.0 ± 11.3	
PPARA rs4823613	AA (n = 155)	-47.4 ± 11.6	
A>G	AG (n = 101)	-47.5 ± 11.4	<i>P</i> >0.05
	GG (n = 14)	-45.0 ± 11.4	
CYP3A enzyme acti	vity —	Downstream activi metabolites of sim	ty vastatin
	Hu M,Tom	inson B. Pharmacogeno	mics (2013) 14(1), 25-34

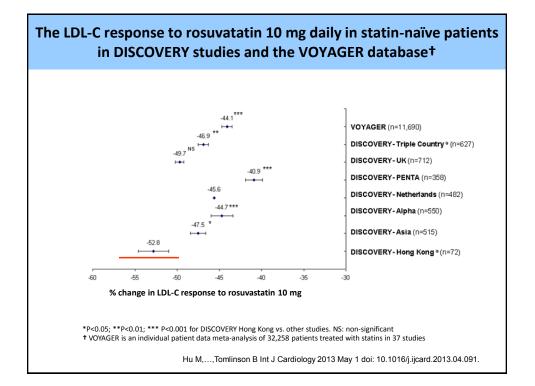


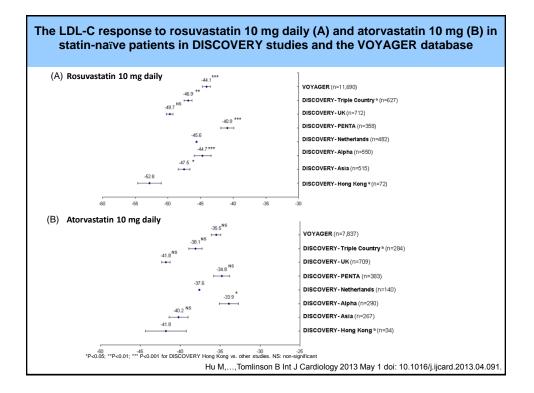


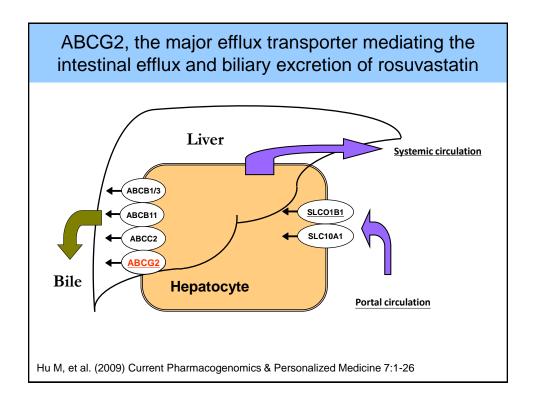


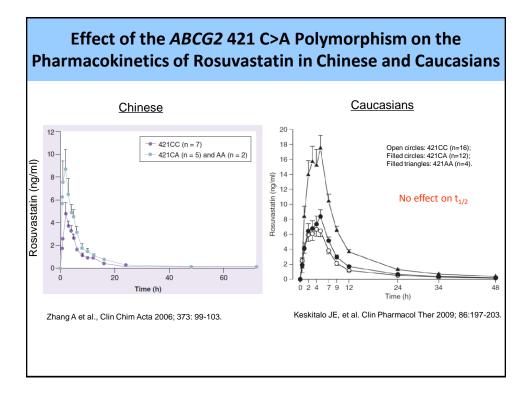


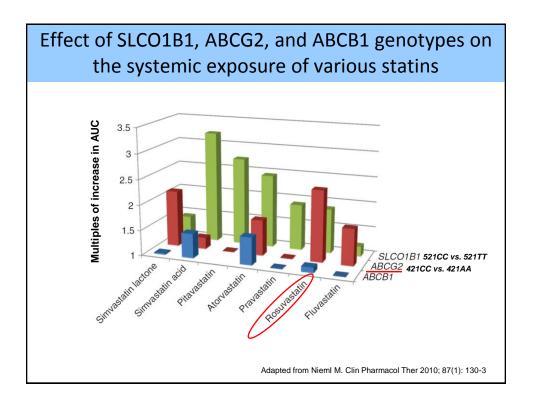
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Letter to the Editor							
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and Caucasians?							
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DISCOVERY Studi	es						
Miao Hu ^a , Sandra S.	H. Lui ^a , Gary	/ T.C. Ko ^b , Brian	Tomlinson ^{a,*}				
* Department of Medicine and Therap	, , ,						
^b Hong Kong Institute of Diabetes and	Obesity, the Chinese Un	iversity of Hong Kong, Hong	Kong SAR				
udy and patient characteristic	s at baseline of the	e various DISCOVERY s	tudies.				
· ·		Asia	Alpha	Netherlands	PENTA	UK	Triple country
	Hong Kong	Maid		recirci unas			
Countries	Hong Kong	China, Hong Kong, Malaysia, Korea, Taiwan, and	93 countries in eastern Europe, Central and South America, and	Netherlands	Brazil, Colombia, Mexico, Portugal, and Venezuela	UK	Finland, Iceland, and Ireland
Countries Randomized population		China, Hong Kong, Malaysia, Korea,	93 countries in eastern Europe, Central and		Brazil, Colombia, Mexico, Portugal,	UK 1847	Finland, Iceland,
	Hong Kong	China, Hong Kong, Malaysia, Korea, Taiwan, and Thailand	93 countries in eastern Europe, Central and South America, and the Middle East	Netherlands	Brazil, Colombia, Mexico, Portugal, and Venezuela		Finland, Iceland, and Ireland
Randomized population	Hong Kong	China, Hong Kong, Malaysia, Korea, Taiwan, and Thailand 1482	93 countries in eastern Europe, Central and South America, and the Middle East 1506	Netherlands 1215 R 10 vs. A 10 vs. S 20 vs. P 40	Brazil, Colombia, Mexico, Portugal, and Venezuela 1124	1847 R 10 vs. A 10 vs. S 20	Finland, Iceland, and Ireland
Randomized population Randomized groups (ratio)	Hong Kong 126 R 10 vs. A 10 (2:1)	China, Hong Kong, Malaysia, Korea, Taiwan, and Thailand 1482 R 10 vs. A 10 (2:1)	93 countries in eastern Europe, Central and South America, and the Middle East 1506 R 10 vs. A 10 (2:1)	Netherlands 1215 R 10 vs. A 10 vs. S 20 vs. P 40 (3:1:1:1)	Brazil, Colombia, Mexico, Portugal, and Venezuela 1124 R 10 vs. A 10 (1:1)	1847 R 10 vs. A 10 vs. S 20 (2:2:1)	Finland, Iceland, and Ireland 1024 R 10 vs. A 10 (2:1)
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Randomized population Randomized groups (ratio) Naïve/switched (%)	Hong Kong 126 R 10 vs. A 10 (2:1) 88/12 R: 57 ± 11.2;	China, Hong Kong, Malaysia, Korea, Taiwan, and Thailand 1482 R 10 vs. A 10 (2:1) 67/33 R: 60 \pm 10.3;	93 countries in eastern Europe, Central and South America, and the Middle East 1506 R 10 vs. A 10 (2:1) 60/40	Netherlands 1215 R 10 vs. A 10 vs. S 20 vs. P 40 (3:1:1:1) 77/22 R: 61 ± 9.7;	Brazil, Colombia, Mexico, Portugal, and Venezuela 1124 R 10 vs. A 10 (1:1) 69/13	1847 R 10 vs. A 10 vs. S 20 (2:2:1) 100/0	Finland, Iceland, and Ireland 1024 R 10 vs. A 10 (2:1) R: 87/13; A: 86/14
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Randomized population Randomized groups (ratio) Naïve/switched (%) Age (years) Male (%)	Hong Kong 126 R 10 vs. A 10 (2:1) 88/12 R: 57 ± 11.2; A: 61 ± 9.6 52	China, Hong Kong, Malaysia, Korea, Taiwan, and Thailand 1482 R 10 vs. A 10 (2:1) 67/33 R: 60 ± 10.3 ; A: 61 ± 10.2 51	93 countries in eastern Europe, Central and South America, and the Middle East 1506 R 10 vs. A 10 (2:1) 60/40 58 \pm 10.8 48	Netherlands 1215 R 10 vs. A 10 vs. S 20 vs. P 40 (3:1:1:1) 77/22 R: 61 ± 9.7; A: 62 ± 9.9 59	Brazil, Colombia, Mexico, Portugal, and Venezuela 1124 R 10 vs. A 10 (1:1) 69/13 59 ± 11.2 41	1847 R 10 vs. A 10 vs. S 20 (2:2:1) 100/0 67 ± 8.7 63	Finland, Iceland, and Ireland 1024 R 10 vs. A 10 (2:1) R: 87/13; A: 86/14 61 ± 10.4 56
Randomized population Randomized groups (ratio) Naïve/switched (%) Age (years) Male (%) CHD (%)	Hong Kong 126 R 10 vs. A 10 (2:1) 88/12 R: 57 ± 11.2; A: 61 ± 9.6 52 25	China, Hong Kong, Malaysia, Korea, Taiwan, and Thailand 1482 R 10 vs. A 10 (2:1) 67/33 R: 60 ± 10.3; A: 61 ± 10.2 51 55	93 countries in eastern Europe, Central and South America, and the Middle East 1506 R 10 vs. A 10 (2:1) 60/40 58 \pm 10.8 48 44	Netherlands 1215 R 10 vs. A 10 vs. S 20 vs. P 40 (3:1:1:1) 77/22 R: 61 ± 9.7; A: 62 ± 9.9 59 27	Brazil, Colombia, Mexico, Portugal, and Venezuela 1124 R 10 vs. A 10 (1:1) 69/13 59 ± 11.2 41 37	1847 R 10 vs. A 10 vs. S 20 (2:2:1) 100/0 67 ± 8.7 63 41	Finland, Icel and, and Ireland 1024 R 10 vs. A 10 (2:1) R: 87/13; A: 86/14 61 ± 10.4 56 28
Randomized population Randomized groups (ratio) Naïve/switched (%) Age (years) Male (%) CHD (%) Diabetes (%)	Hong Kong 126 R 10 vs. A 10 (2:1) 88/12 R: 57 ± 11.2; A: 61 ± 9.6 52 25 74	$\begin{array}{l} {\rm China, Hong Kong, \\ {\rm Malaysia, Korea, } \\ {\rm Taiwan, and } \\ {\rm Taiwan, and } \\ {\rm Taiwan, and } \\ {\rm R 10 vs. A 10 } \\ {\rm (2:1)} \\ {\rm 67/33} \\ {\rm R: 60 \pm 10.3; } \\ {\rm A: 61 \pm 10.2 } \\ {\rm 51} \\ {\rm 55} \\ {\rm 45} \end{array}$	93 countries in eastern Europe, Central and South America, and the Middle East 1506 R 10 vs. A 10 (2:1) 60/40 58 \pm 10.8 48 44 35	Netherlands 1215 R 10 vs. A 10 vs. S 20 vs. P 40 (3:1:1:1) 77/22 R: 61 ± 9.7; A: 62 ± 9.9 59 27 26	Brazil, Colombia, Mexico, Portugal, and Venezuela 1124 R 10 vs. A 10 (1:1) 69/13 59 ± 11.2 41 37 19	1847 R 10 vs. A 10 vs. S 20 (2:2:1) 100/0 67 ± 8.7 63 41 17	Finland, Iceland, and Ireland 1024 R 10 vs. A 10 (2:1) R: $87/13$; A: $86/14$ 61 ± 10.4 56 28 27
Randomized population Randomized groups (ratio) Naīve/switched (%) Age (years) Male (%) CHD (%) Diabetes (%) Baseline LDL-c in all study subjects (mmol/L)	Hong Kong 126 R 10 vs. A 10 (2:1) 88/12 R: 57 ± 11.2; A: 61 ± 9.6 52 25 74 R: 4.4 ± 0.7; A: 4.5 ± 1.0	$\begin{array}{l} {\rm China, Hong Kong, \\ {\rm Malayai, Korea, \\ {\rm Taiwan, and \\ {\rm Thailand} \\ {\rm 1482} \\ {\rm R} : 10{\rm vs.}{\rm A} : 10 \\ {\rm (2:1)} \\ {\rm 67/33} \\ {\rm R} : 60\pm10.3; \\ {\rm A} : 61\pm10.2 \\ {\rm 515} \\ {\rm 55} \\ {\rm 45} \\ {\rm R} : {\rm A} : 1\pm0.8; \\ {\rm A} : {\rm A2}\pm0.8 \end{array}$	$\begin{array}{c} 93 \ countries in eastern \\ Europe, Central and \\ South America, and \\ the Middle East \\ 1506 \\ R \ 10 \ vs. A \ 10 \\ (2:1) \\ 60/40 \\ 58 \ \pm \ 10.8 \\ 48 \\ 44 \\ 35 \\ R: 46 \ (N); \ 4.1 \ (S) \\ A: 4.6 \ (N); 4.1 \ (S) \end{array}$	$\begin{array}{c} \text{Netherlands} \\ \hline 1215 \\ \text{R 10 vs, A 10 vs, } \\ \text{S 20 vs, P 40} \\ (3;1;1;1) \\ 77/22 \\ \text{R: } 61 \pm 9.7; \\ \text{A: } 62 \pm 9.9 \\ 59 \\ 59 \\ 27 \\ 26 \\ \text{R: } 4.5 \pm 0.8; \\ \text{A: } 4.4 \pm 0.7 \\ \end{array}$	$\begin{array}{l} Brazil, Colombia, \\ Mexico, Portugal, \\ and Venezuela \\ 1124 \\ R 10 vs. A 10 \\ (1:1) \\ 69/13 \\ 59 \pm 11.2 \\ 41 \\ 37 \\ 19 \\ 4.4 \pm 1.0 \end{array}$	$\begin{array}{c} 1847 \\ R \ 10 \ vs. \ A \ 10 \\ vs. \ 5 \ 20 \\ (2:2:1) \\ 100/0 \\ 67 \ \pm \ 8.7 \\ 63 \\ 41 \\ 17 \\ 4.5 \ \pm \ 0.7 \end{array}$	$\label{eq:response} \begin{array}{l} Finland, Iceland, \\ and Ireland \\ \hline 1024 \\ R \ 10 \ vs. \ A \ 10 \\ (2:1) \\ R: \ 87/13; \\ A: \ 86/14 \\ 61 \ \pm \ 10.4 \\ \hline 56 \\ 28 \\ 27 \\ R: \ 4.5 \ \pm \ 0.8; \end{array}$
Randomized population Randomized groups (ratio) Naïve/switched (%) Age (years) Male (%) CHD (%) Diabetes (%) Baseline LD-C in all study	Hong Kong 126 R 10 vs. A 10 (2:1) 88/12 R: 57 ± 11.2; A: 61 ± 9.6 52 25 74 R: 4.4 ± 0.7; A: 4.5 ± 1.0	$\begin{array}{l} {\rm China, Hong Kong, \\ {\rm Malayai, Korea, \\ {\rm Taiwan, and \\ {\rm Thailand} \\ {\rm 1482} \\ {\rm R} : 10{\rm vs.}{\rm A} : 10 \\ {\rm (2:1)} \\ {\rm 67/33} \\ {\rm R} : 60\pm10.3; \\ {\rm A} : 61\pm10.2 \\ {\rm 515} \\ {\rm 55} \\ {\rm 45} \\ {\rm R} : {\rm A} : 1\pm0.8; \\ {\rm A} : {\rm A2}\pm0.8 \end{array}$	$\begin{array}{c} 93 \ countries in eastern \\ Europe, Central and \\ South America, and \\ the Middle East \\ 1506 \\ R \ 10 \ vs. A \ 10 \\ (2:1) \\ 60/40 \\ 58 \ \pm \ 10.8 \\ 48 \\ 44 \\ 35 \\ R: 46 \ (N); \ 4.1 \ (S) \\ A: 4.6 \ (N); 4.1 \ (S) \end{array}$	$\begin{array}{c} \text{Netherlands} \\ \hline 1215 \\ \text{R 10 vs, A 10 vs, } \\ \text{S 20 vs, P 40} \\ (3;1;1;1) \\ 77/22 \\ \text{R: } 61 \pm 9.7; \\ \text{A: } 62 \pm 9.9 \\ 59 \\ 59 \\ 27 \\ 26 \\ \text{R: } 4.5 \pm 0.8; \\ \text{A: } 4.4 \pm 0.7 \\ \end{array}$	$\begin{array}{l} Brazil, Colombia, \\ Mexico, Portugal, \\ and Venezuela \\ 1124 \\ R 10 vs. A 10 \\ (1:1) \\ 69/13 \\ 59 \pm 11.2 \\ 41 \\ 37 \\ 19 \\ 4.4 \pm 1.0 \end{array}$	$\begin{array}{c} 1847 \\ R \ 10 \ vs. \ A \ 10 \\ vs. \ 5 \ 20 \\ (2:2:1) \\ 100/0 \\ 67 \ \pm \ 8.7 \\ 63 \\ 41 \\ 17 \\ 4.5 \ \pm \ 0.7 \end{array}$	$\label{eq:response} \begin{array}{l} Finland, Iceland, \\ and Ireland \\ \hline 1024 \\ R \ 10 \ vs. \ A \ 10 \\ (2:1) \\ R: \ 87/13; \\ A: \ 86/14 \\ 61 \ \pm \ 10.4 \\ \hline 56 \\ 28 \\ 27 \\ R: \ 4.5 \ \pm \ 0.8; \end{array}$
Randomized population Randomized groups (ratio) Naīve/switched (%) Age (years) Male (%) CHD (%) Diabetes (%) Baseline LDL-c in all study subjects (mmol/L)	Hong Kong 126 R 10 vs. A 10 (2:1) 88/12 R: 57 ± 11.2; A: 61 ± 9.6 52 25 74 R: 4.4 ± 0.7; A: 4.5 ± 1.0	$\begin{array}{l} {\rm China, Hong Kong, \\ {\rm Malayai, Korea, \\ {\rm Taiwan, and \\ {\rm Thailand} \\ {\rm 1482} \\ {\rm R} : 10{\rm vs.}{\rm A} : 10 \\ {\rm (2:1)} \\ {\rm 67/33} \\ {\rm R} : 60\pm10.3; \\ {\rm A} : 61\pm10.2 \\ {\rm 515} \\ {\rm 55} \\ {\rm 45} \\ {\rm R} : {\rm A} : 1\pm0.8; \\ {\rm A} : {\rm A2}\pm0.8 \end{array}$	$\begin{array}{c} 93 \ countries in eastern \\ Europe, Central and \\ South America, and \\ the Middle East \\ 1506 \\ R \ 10 \ vs. A \ 10 \\ (2:1) \\ 60/40 \\ 58 \ \pm \ 10.8 \\ 48 \\ 44 \\ 35 \\ R: 46 \ (N); \ 4.1 \ (S) \\ A: 4.6 \ (N); 4.1 \ (S) \end{array}$	$\begin{array}{c} \text{Netherlands} \\ \hline 1215 \\ \text{R 10 vs, A 10 vs, } \\ \text{S 20 vs, P 40} \\ (3;1;1;1) \\ 77/22 \\ \text{R: } 61 \pm 9.7; \\ \text{A: } 62 \pm 9.9 \\ 59 \\ 59 \\ 27 \\ 26 \\ \text{R: } 4.5 \pm 0.8; \\ \text{A: } 4.4 \pm 0.7 \\ \end{array}$	$\begin{array}{l} Brazil, Colombia, \\ Mexico, Portugal, \\ and Venezuela \\ 1124 \\ R 10 vs. A 10 \\ (1:1) \\ 69/13 \\ 59 \pm 11.2 \\ 41 \\ 37 \\ 19 \\ 4.4 \pm 1.0 \end{array}$	$\begin{array}{c} 1847 \\ R \ 10 \ vs. \ A \ 10 \\ vs. \ 5 \ 20 \\ (2:2:1) \\ 100/0 \\ 67 \ \pm \ 8.7 \\ 63 \\ 41 \\ 17 \\ 4.5 \ \pm \ 0.7 \end{array}$	

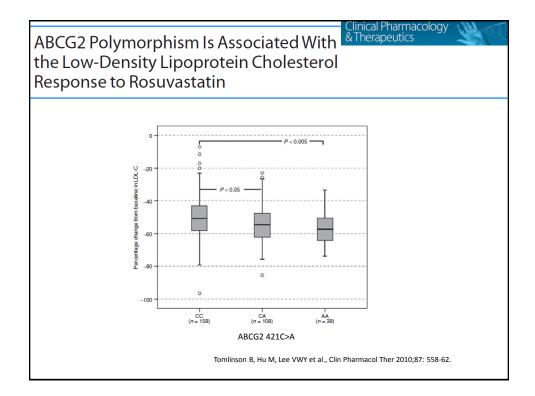


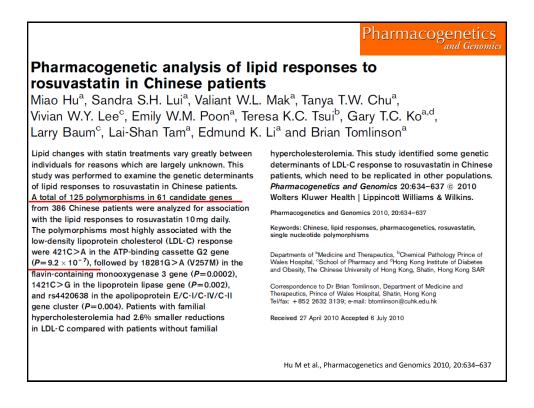


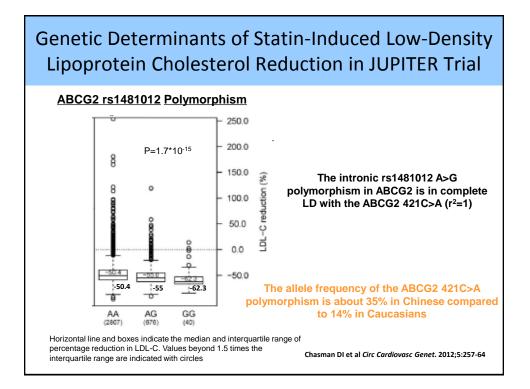






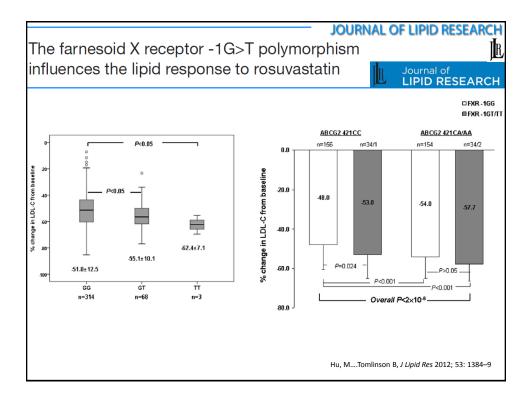


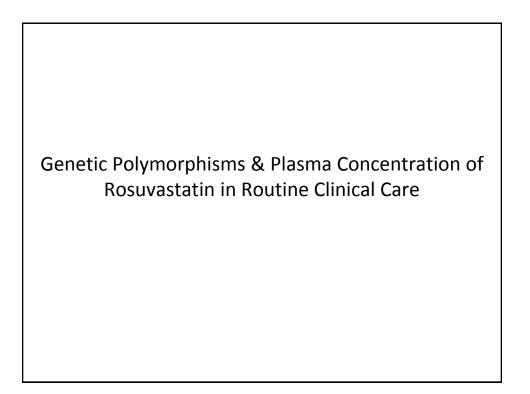


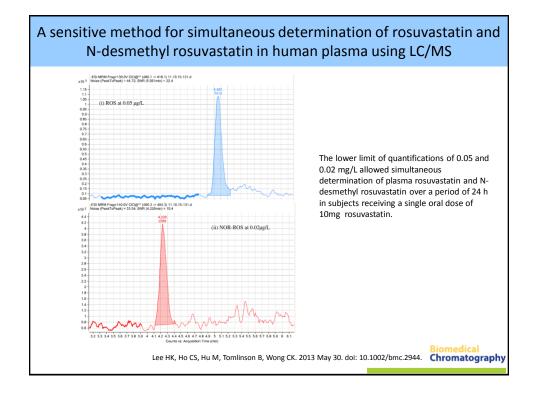


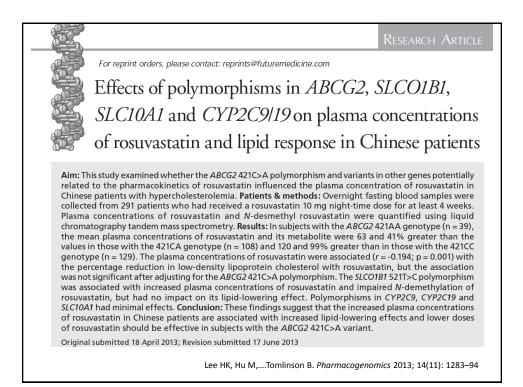
Farnesoid X receptor (FXR), a bile-acid-activated nuclear receptor

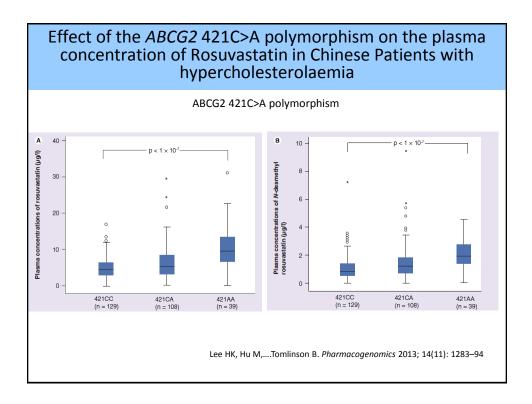
- FXR is a nuclear receptor and a receptor for bile acids
- Activation of FXR leads to altered expression of many genes responsible for bile acid and lipid and glucose metabolism and transport
- FXR also regulates multiple drug metabolizing enzymes and drug transporters (e.g. SLCO1B1, SLCO1B3, NTCP and some efflux transporters) by binding to FXR response elements and promoting transcription of target genes

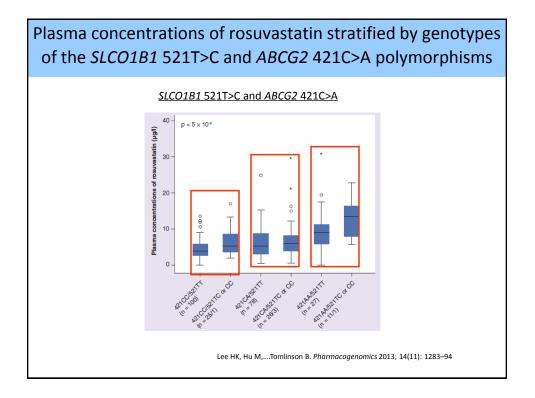












Association between genotypic and phenotypic factors, and the plasma concentrations of rosuvastatin

Variables	Model 1		
	β	p-value	R ²
ABCG2 421C>A polymorphism (1 = CC, 2 = CA, 3 = AA)	0.379	<1 × 10 ⁻¹⁰	0.145
Bodyweight (kg)	-0.238	<1 × 10 ⁻⁴	0.043
Hypertension $(0 = no, 1 = yes)$	0.192	<0.001	0.039
SLCO1B1 521T>C polymorphism (1 = TT, 2 = TC or CC)	0.123	0.024	0.015

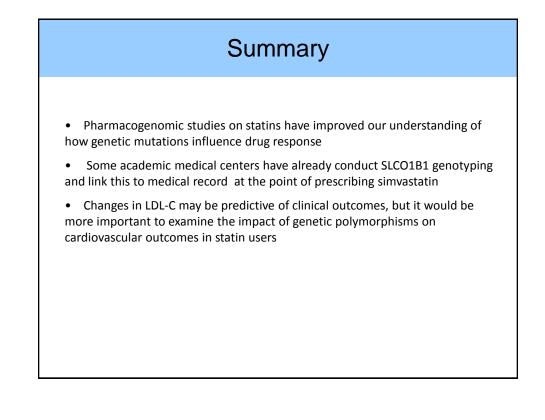
Lee HK, Hu M,....Tomlinson B. Pharmacogenomics 2013; 14(11): 1283–94

Genome-wide association studies on the pharmacogenomics of statins

Statins	Study population	Reductions in LDL-C or other cardiovascular risk factors [†]	Myopathy
Atorvastatin	TNT	None, but APOE significant at genome-wide level and PCSK9, HMGCR significant in candidate gene analysis	NA
Atorvastatin	CARDS, ASCOT and PROSPER	APOE and LPA	NA
Pravastatin	CARE, WOSCOPS and PROSPER/PHASE	NA	NA
Rosuvastatin	JUPITER	<u>ABCG2</u> , APOE, LPA (PCSK9 gene-significant);	NA
		None for changes in CRP ⁺	NA
		ABCG2 and LPA for changes in Lp-PLA ₂ , but no association after adjustment for LDL-C changes [†]	NA
Simvastatin	HPS	None, but <i>LPA</i> and <i>APOE</i> signficant in candidate gene analysis	NA
Simvastatin	SEARCH	NA	SLCO1B1

Summary

- The hepatic uptake transporter SLCO1B1 plays an important role in determining the plasma concentrations and the risk of myopathy of statins
- The functional SNP 521 T>C in *SLCO1B1* significantly increases the pharmacokinetics of statins and increases the risk of myopathy of simvastatin, but it had little effect on the hepatic exposure and lipid-lowering effect of statins
- The efflux transporter ABCG2 appears to significantly affect the absorption and the biliary excretion of the substrate statins.
- The loss-of-function mutation 421C>A in *ABCG2* is the major genetic determinant of the pharmacokinetics and the lipid-lowering effect of rosuvastatin.



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Thank You For Your Attention!