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Genetic and seasonal determinants of vitamin D status in Confederated Salish and Kootenai Tribes (CSKT) participants

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Staples, Jack W.; George, Kathleen M.; Dalton, Rachel; Ellerbeck, Erin E.; Krause, Genevieve; Lietch, Tianna; Aliwarga, Theresa; McDonald, Matthew G.; Phillips, Brian; Muzquiz, LeeAnna I.; Nickerson, Deborah A.; Thornton, Timothy A.; Thummel, Kenneth E.; and Woodahl, Erica L., "Genetic and seasonal determinants of vitamin D status in Confederated Salish and Kootenai Tribes (CSKT) participants" (2022). *UM Graduate Student Research Conference (GradCon)*. 6. https://scholarworks.umt.edu/gsrc/2022/posters/6

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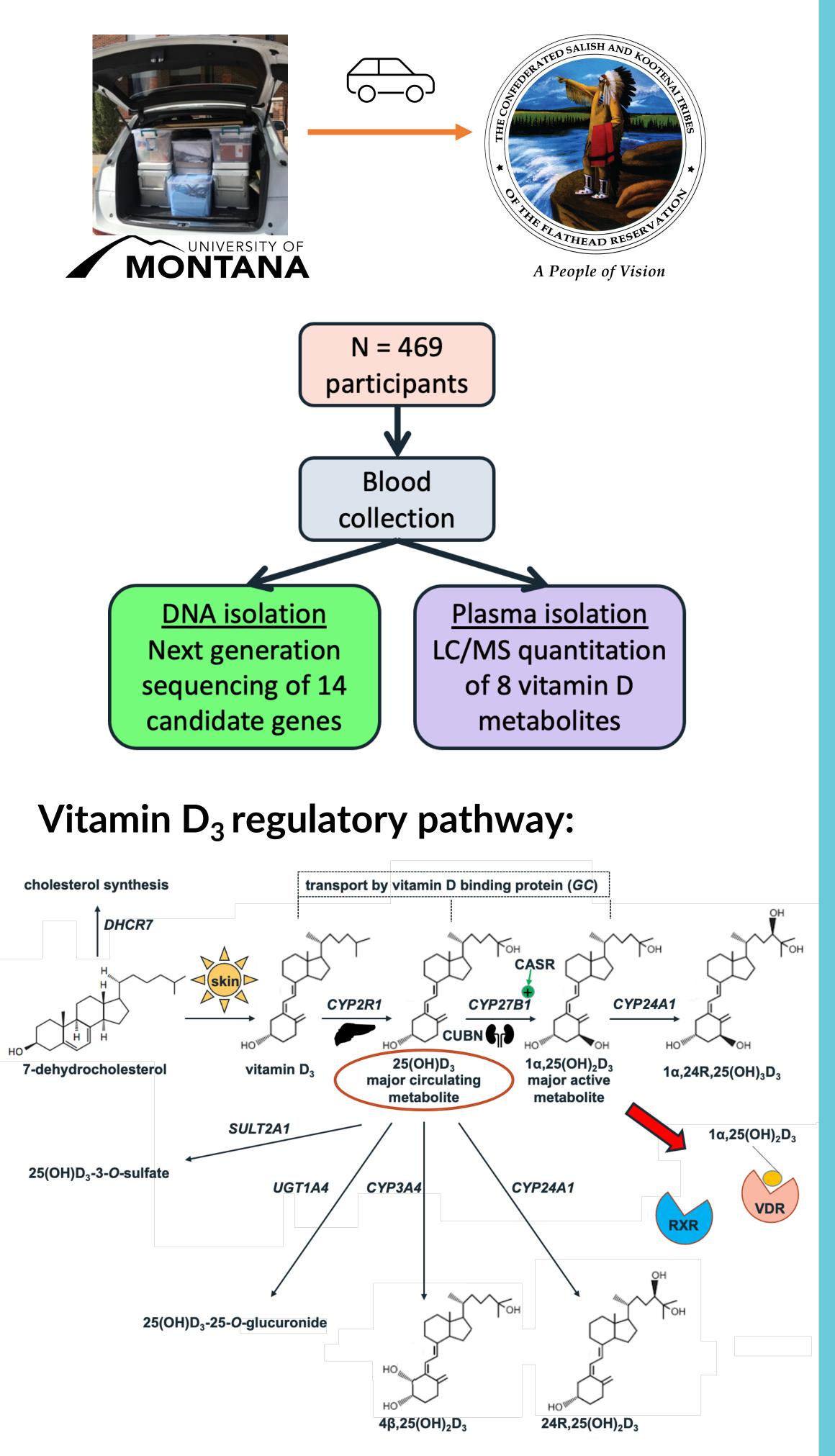
Genetic and seasonal determinants of vitamin D status in Confederated Salish and Kootenai Tribes (CSKT) participants



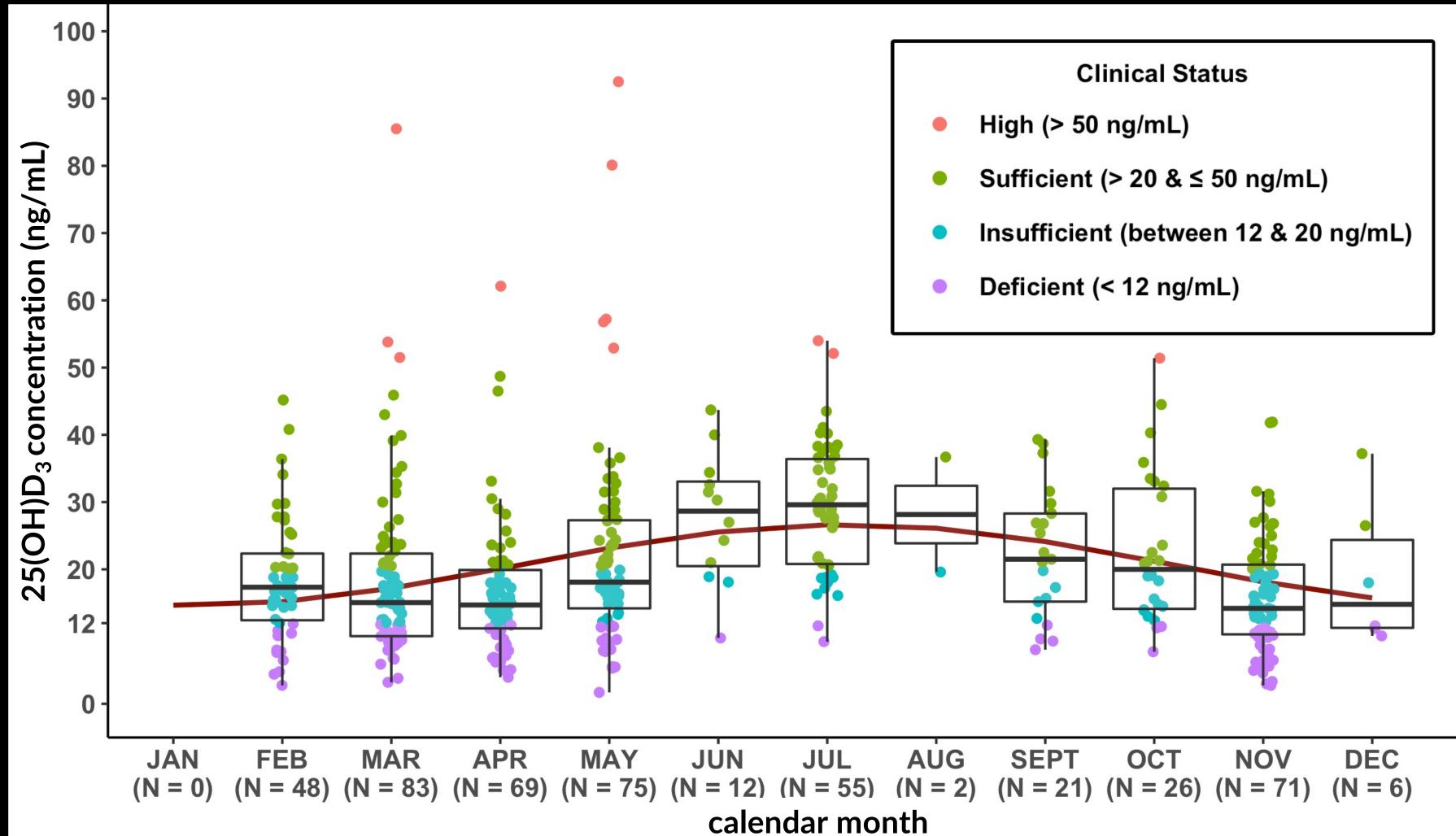
PRESENTER: Jack Staples jack.staples@umontana.edu

Background: Vitamin D levels vary globally in human populations due to genetics, geography, and other demographic factors (e.g., age, BMI, and gender). To improve our understanding of contributors to vitamin D levels, we conducted a **candidate-gene** study in partnership with the CSKT.

Methods: Our team traveled over 5,000 total miles to conduct this study.

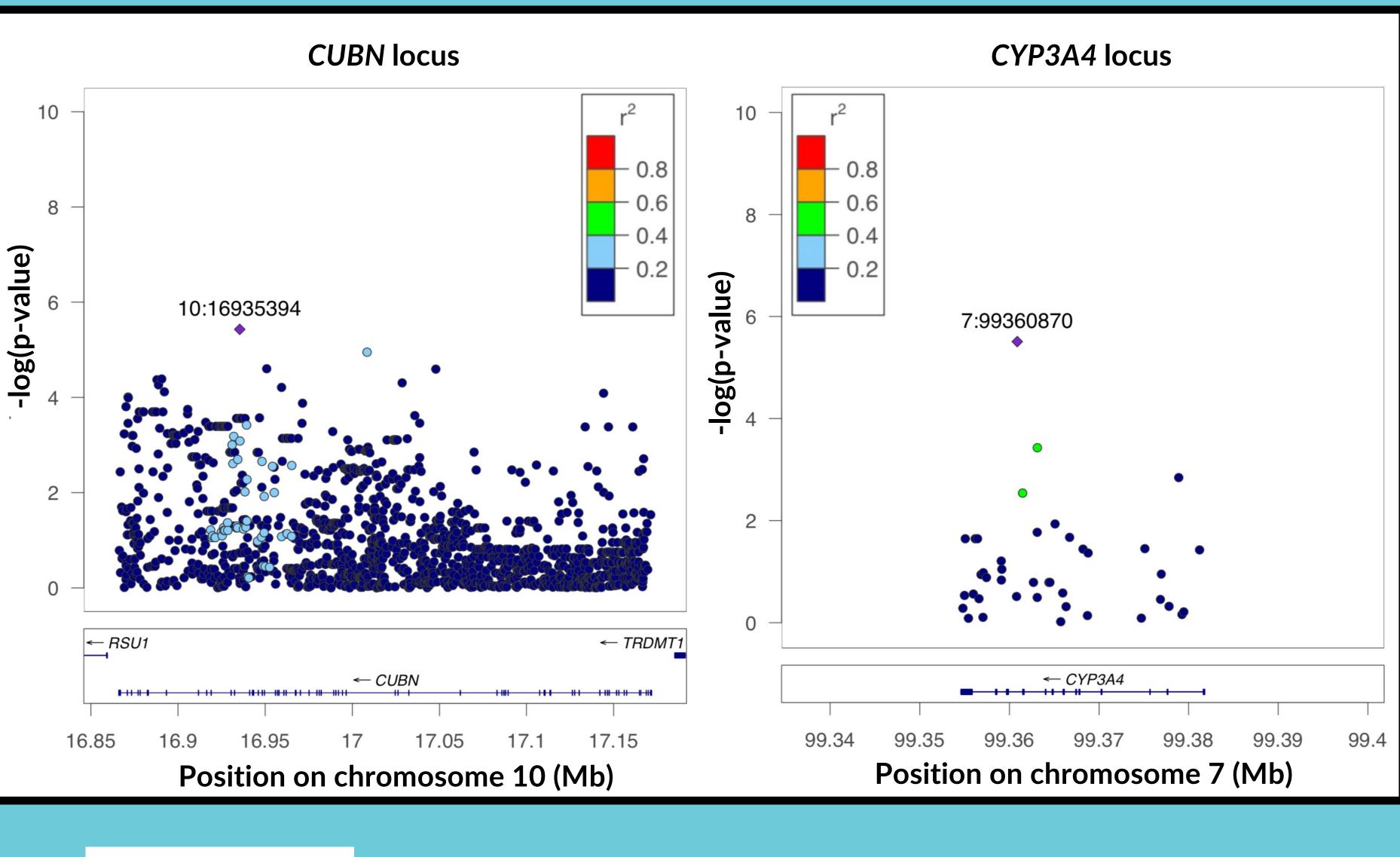


55% of CSKT had vitamin D levels below sufficiency (< 20 ng/mL)

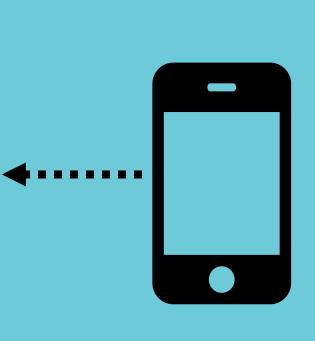


4 variants in CUBN and 1 variant in **CYP3A4** were significantly associated

with 25(OH)D₃ concentration





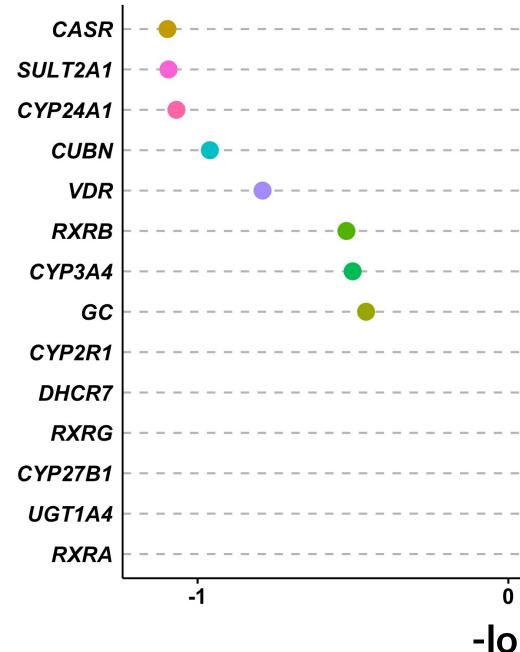


Take a picture to visit the **Precision Medicine Project website!**

Multivariate regression: 23% of 25(OH)D₃ variability explained

Variant or covariate	Beta ± S.E.	P-value
rs59819645 (CUBN)	-3.94 ± 1.08	0.000307***
rs7089377 (CUBN)	-2.42 ± 0.696	0.00056***
rs34859798 (CUBN)	1.72 ± 0.951	0.0713
rs77520345 (CUBN)	7.34 ± 2.54	0.00398**
rs4646440 (CUBN)	-3.28 ± 0.728	8.33e-06***
Gender = Male	-2.01 ± 0.966	0.0385*
Age	0.104 ± 0.0391	0.00844**
BMI	-0.239 ± 0.0734	0.00125**
Season = SeptNov.	-8.5 ± 1.48	1.63e-08***
Season = DecFeb.	-10.3 ± 1.7	3.24e-09***
Season = MarMay	-9.72 ± 1.44	4.63e-11***
*p < 0.05, *p < 0.0	1, *p < 0.001	

sufficiency [25(OH)D < 20 ng/mL]:



Acknowledgements:

We thank the CSKT research participants who graciously volunteered their time. This work is supported by the Northwest-Alaska Pharmacogenomics Research Network (P01GM116691) and the American Indian and Alaska Native Clinical and Translational Research Program (U54GM115371).



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Pooled rare variant analysis of 25(OH)D significant associations in UGT1A and RXRA

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-log(p-value)

