

Defining and Measuring Safety of Combinatorial Pharmacogenomic Testing for Patients with Major Depressive Disorder (MDD)

Sagar V. Parikh, MD¹; Gabriela K. Khazanov, PhD²; Michael E. Thase, MD²; Anthony J. Rothschild, MD³; Boadie W. Dunlop, MD⁴; Charles DeBattista, DMH, MD⁵; Charles R. Conway, MD⁶; Brent P. Forester, MD, MSc⁷; Richard C. Shelton, MD⁸; Matthew Macaluso, DO⁹; James Li, MS⁹; Kunbo Yu, MS⁹; Stephanie Meek, PhD¹⁰; Michael R. Jablonski, PhD⁹; John F. Greden, MD¹

OBJECTIVE

- Pharmacogenomic (PGx) tests are increasingly used to guide medication prescribing in MDD. While efficacy of PGx is promising, the potential for patient harm should be assessed. Here, we use data from the GUIDED trial to evaluate the safety of using the GeneSight test to guide treatment decisions, looking specifically for evidence of patient harms after medications changes are made.

METHODS

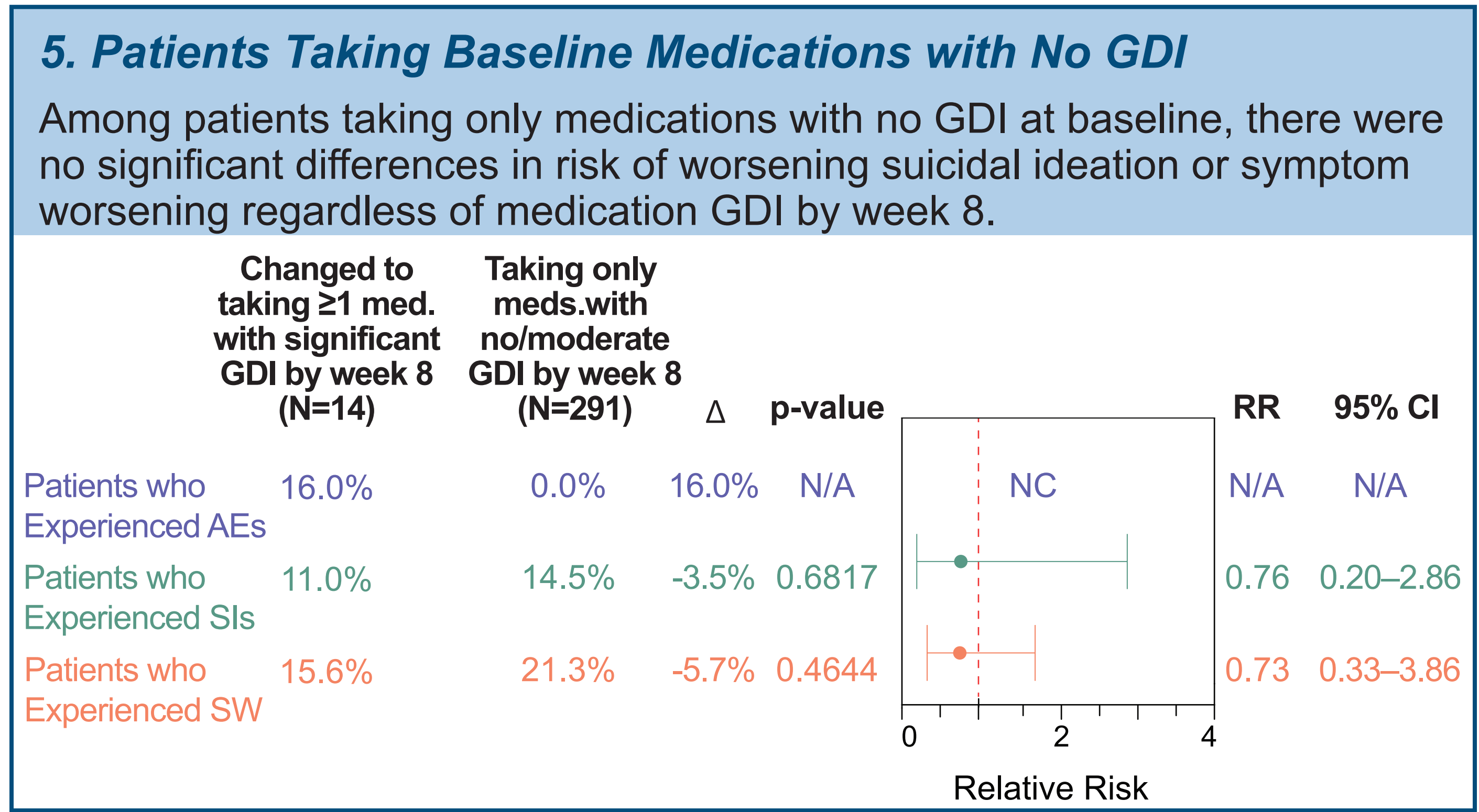
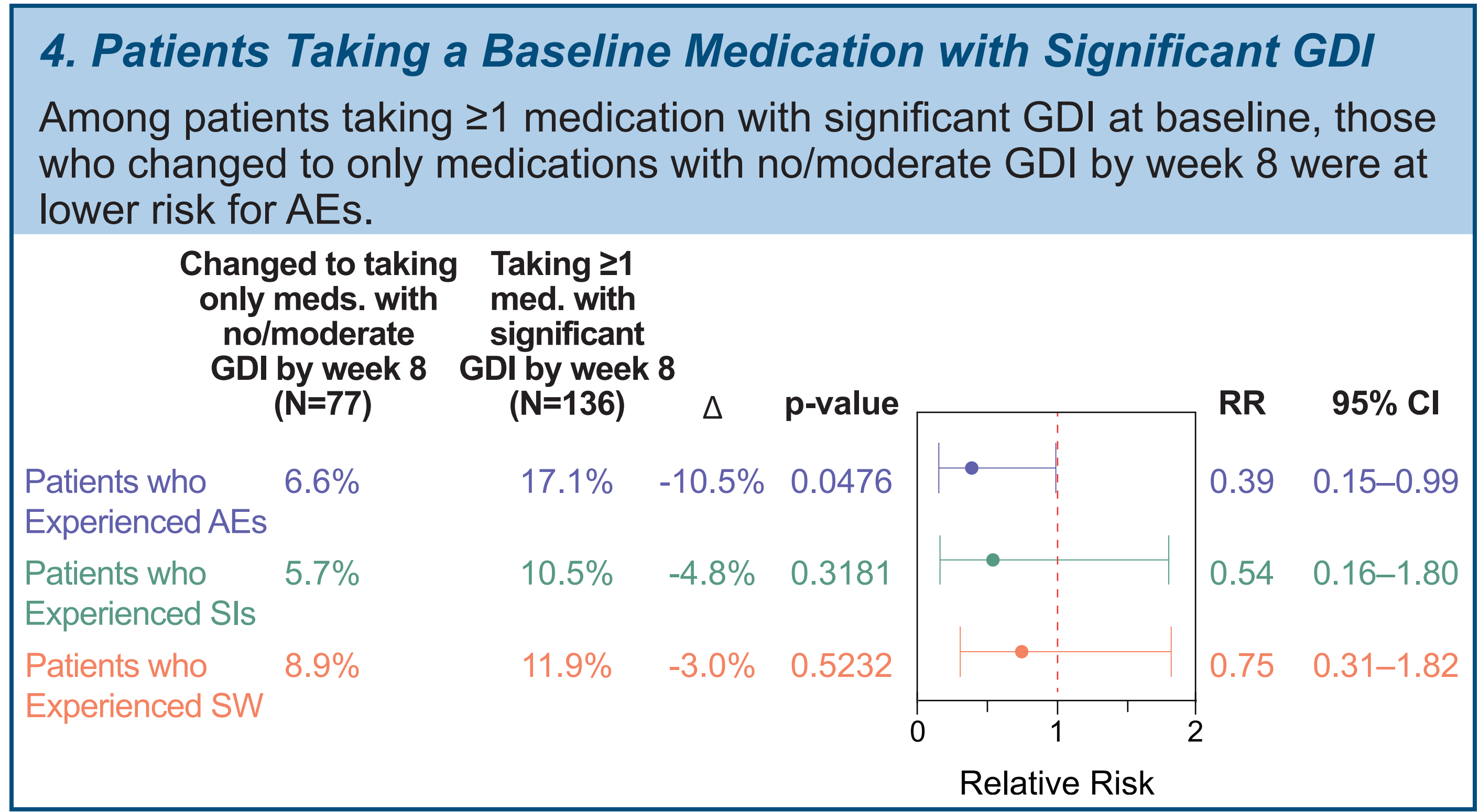
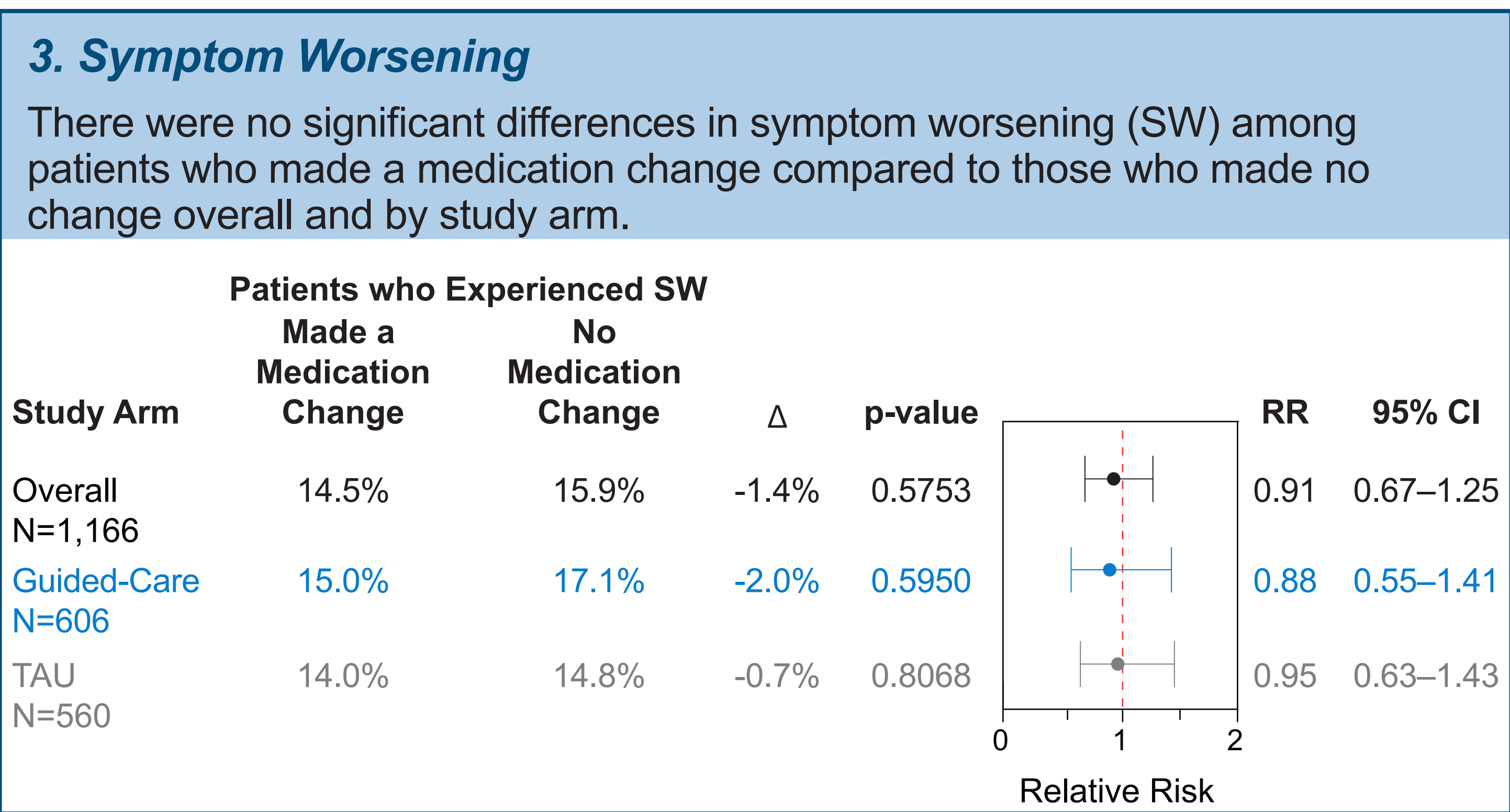
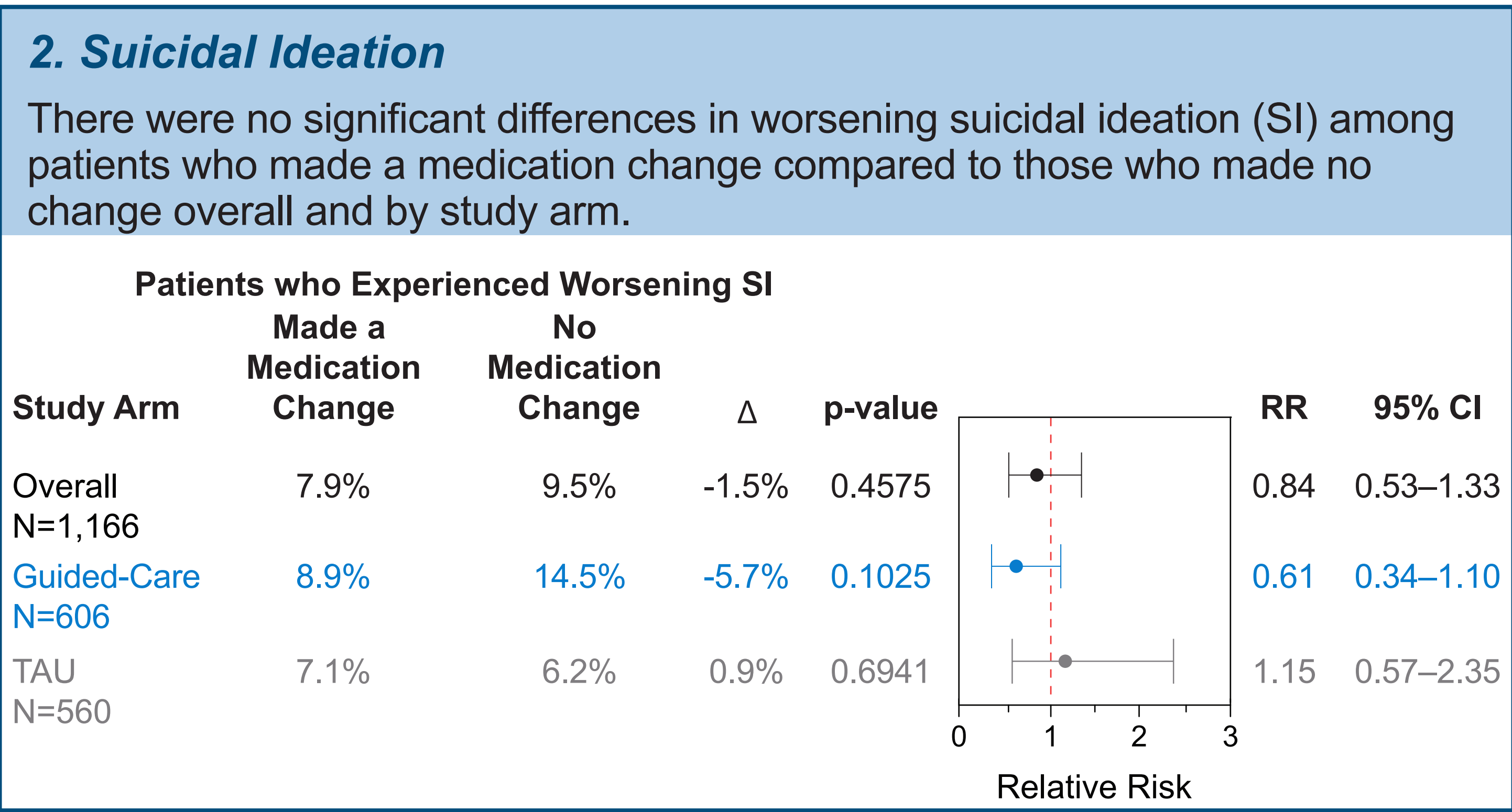
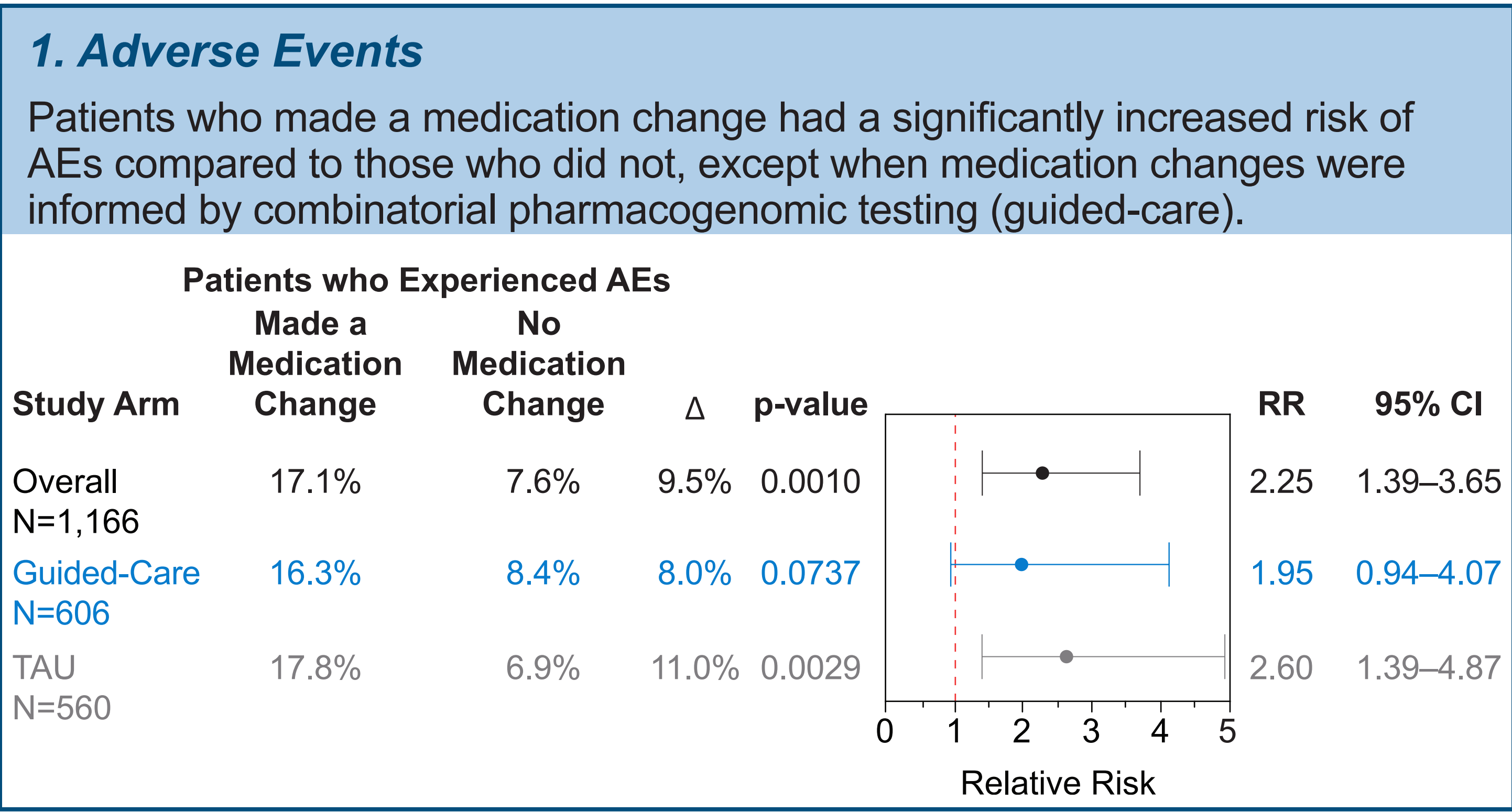
COHORT

- GUIDED was a patient- and rater-blinded, randomized, controlled trial including patients diagnosed with MDD who had an inadequate response to ≥1 psychotropic medication.
- All patients received combinatorial pharmacogenomic testing and medications were categorized according to the level of predicted gene-drug interactions (no, moderate, or significant GDI).
- Patients were randomized 1:1 to the combinatorial pharmacogenomic guided-care arm or treatment as usual (TAU). Patients and raters were blinded through week 8. Clinicians were blinded to pharmacogenomic test results for patients in TAU.

ANALYSIS

- Patient harms were defined as:
 - Adverse Events** (AEs, present/absent)
 - Worsening Suicidal Ideation** (increase ≥1 on the HAM-D question)
 - Symptom Worsening** (HAM-D17 increase of ≥1).
- The relative risk of each measure was assessed for patients who changed medications [add and/or drop a medication] and those who made no change.
- Relative risk was also assessed according to medication GDI at baseline and week 8.
 - Relative risk >1 indicates higher risk among patients who made a medication change.

Presented at NEI on November 5-8, 2020



CONCLUSION

- There was no increased patient harm when combinatorial pharmacogenomic testing was used to inform treatment decisions.
- For patients with significant GDI, patient safety may be improved when treatment decisions align with the combinatorial pharmacogenomic test results.
- This indicates that combinatorial pharmacogenomic testing can be adopted safely into clinical practice without increasing the risk for adverse clinical outcomes.

AFFILIATIONS

1. University of Michigan Comprehensive Depression Center and Department of Psychiatry and NNDC, Ann Arbor, MI; 2. Perelman School of Medicine of the University of Pennsylvania and the Corporal Michael Crescenz VAMC, Philadelphia, PA; 3. University of Massachusetts Medical School and UMass Memorial Healthcare, Worcester, MA; 4. Emory University School of Medicine, Atlanta, GA; 5. Stanford University School of Medicine, Stanford, CA; 6. Washington University School of Medicine, St. Louis, MO; 7. McLean Hospital, Belmont, MA; 8. The University of Alabama at Birmingham, Birmingham, AL; 9. Myriad Neuroscience, Mason, OH; 10. Myriad Genetics, Inc., Salt Lake City, UT