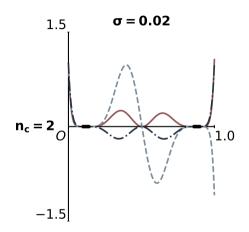
Sharpened CG Iteration Bound for High-contrast Heterogeneous Scalar Elliptic PDEs

Going Beyond Condition Number

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- Linear system of equations Au = b
- We can compute approximations $\mathbf{u}_1, \mathbf{u}_2, \dots$ efficiently using the CG method.
- Main R.Q.: "How can we improve existing estimates on the total number of necessary CG iterations?"



Structure

- Introducing CG
- How Does CG Converge? The Role of Eigenvalues
- Preconditioning: Taming High-Contrast Problems
- High-contrast coefficients: split eigenspectrum
- Towards Sharper Iteration Bounds
- Multi-Cluster Spectra
- How Sharp Are the New Bounds?
- New Bounds in Practice: Using Ritz Values
- Conclusion: Key Takeaways & Future Directions



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- However m_1 can be too pessimistic for high-contrast problems. That is, $m \ll m_1$
- Restate R.Q.: "How do we improve/sharpen the classical bound m_1 for model problem?"



The Role of Eigenvalues

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- Animation: cg_residual_poly
 - Loop through different randomized, clustered spectra, showing r_m for each.
 - Show best and worst case scenario's for CG convergence



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Taming High-Contrast Problems

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- However, on a sample problem PCG with M₁, M₂ needed significantly less iterations than PCG with M₃.
- Restate R.Q.: "How can we construct a CG iteration bound that can distinguish between different preconditioners with similar conditioner numbers?"



The Two-clusters case

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- Note: The transition from classical bounds to multi-cluster bounds could be made more explicit—perhaps with a summary slide or visual.



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- Idea: simple, we split at the largest (relative) gap between subsequent eigenvalues, check if the two resulting clusters would give a sharper bound than just one uniform cluster, that is $m_2(\kappa, \kappa_I, \kappa_r) < m_1(\kappa)$. If so, we split the clusters and repeat the process for each created cluster. If not, we stop partitioning. In the process we keep track of all the extremal eigenvalues. After partitioning, we calculate $m_{N_{\text{cluster}}}$.



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- Restate R.Q.: "How much sharper is $m_{N_{\text{cluster}}}$ compared to $m_1(\kappa)$ for our model problem?"



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 For comparison, m₁ is anywhere from 100 to 1000 times bigger than m, where the difference increases for spectra with larger spectral gaps
- Restate R.Q.: "How can we compute $m_{N_{\text{cluster}}}$ for our model problem in practice?"



Using Ritz Values

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- Animation: ritz_value_migration



