

GitHub Repo

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Contact

Clenshaw Graph Neural Networks

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Model ClenshawGCN

A new **negative** second-order residual.

$$\mathbf{H}^{(\ell+1)} = \sigma((2\tilde{\mathbf{P}}\mathbf{H}^{(\ell)} - \mathbf{H}^{(\ell-1)} + \alpha_\ell \mathbf{H}^*)((1 - \theta_\ell)\mathbf{W}^{(\ell)} + \theta_\ell \mathbf{I}))$$

- Two shortcuts.
- Allows for simulating **arbitrary** polynomial filter upon **Chebyshev basis** (the second kind).
- The **convolution process** mimics **Clenshaw's algorithm**.
- Inherits strengths from both spatial GNNs and spectral GNNs.

Clenshaw's Algorithm: Evaluates the value of $p(x)$ at x_0 where $p(x) = a_0 + a_1 U_1(x) + \dots + a_n U_n(x)$, and U_k is the k -th **Chebyshev polynomial** (the 2nd kind).

Unfolding Process of ClenshawGCN

$$\begin{aligned} b_{n+2}(x_0) &:= 0 \\ b_{n+1}(x_0) &:= 0 \\ b_k(x_0) &:= a_k + 2x_0 b_{k+1}(x_0) - b_{k+2}(x_0) \\ &\quad (k = n, n-1, \dots, 0) \end{aligned}$$

$$\begin{aligned} \mathbf{H}^{(-2)} &= \mathbf{0} \\ \mathbf{H}^{(-1)} &= \mathbf{0} \\ \mathbf{H}^{(\ell+1)} &= 2\tilde{\mathbf{P}}\mathbf{H}^{(\ell)} - \mathbf{H}^{(\ell-1)} + \alpha_\ell \mathbf{H}^{(0)} \\ &\quad (i = 0, 1, \dots, K) \end{aligned}$$

Background

Spatial GNNs

$$\mathbf{H} \mapsto \tilde{\mathbf{P}}\mathbf{H}\mathbf{W}^{(\ell)}$$

Residual connections equipped spatial GNNs

- **Categorization:** Vanilla/initial/dense residuals.
- **Motivations:** Tackling model degradation; enhance expressiveness.

Spectral GNNs

$$\mathbf{x} \mapsto \mathbf{U} \cdot \text{diag}\{\theta_1, \dots, \theta_N\} \cdot \mathbf{U}^T \cdot \mathbf{x}$$

See Footnote.

Polynomial Filters

$$\mathbf{x} \mapsto \mathbf{U} \cdot \text{diag}\{h(\lambda_1), \dots, h(\lambda_N)\} \cdot \mathbf{U}^T \cdot \mathbf{x}$$

$$\mathbf{x} \mapsto h(\tilde{\mathbf{P}})\mathbf{x} \text{ or } \mathbf{b}(\tilde{\mathbf{P}})\mathbf{x}$$

Computed Localized!

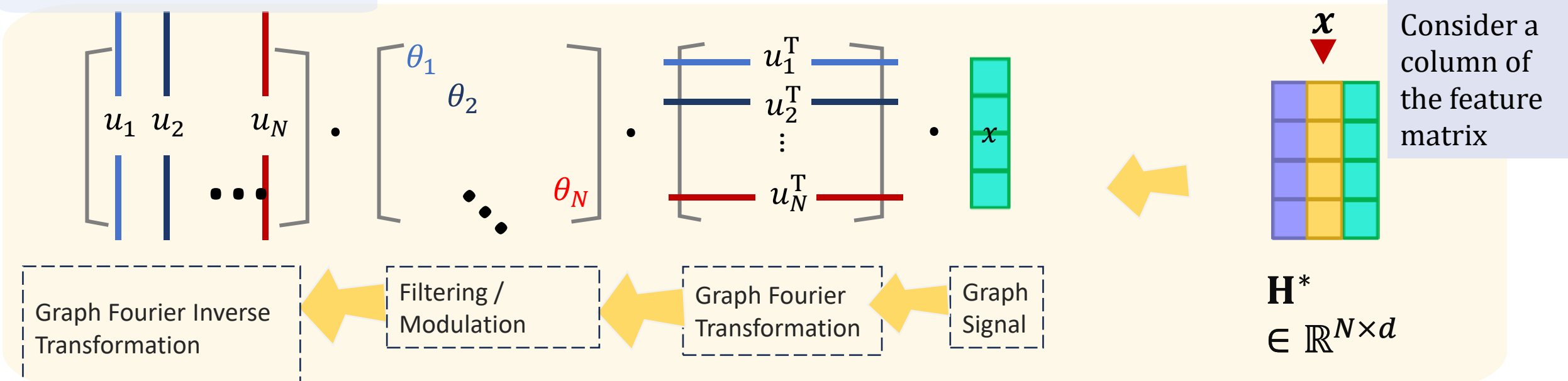
$$\sum_{k=0}^K \alpha_k g_k(\tilde{\mathbf{P}})$$

Feature I:
Arbitrary coefficients.
Feature II:
Utilizing polynomial basis.

Our contribution

- Simple & novel residuals;
- Equip a spatial GCN to simulate a full-featured polynomial filter.
- Gap between spatial & spectral GNNs.

Footnote: on spectral GNN



Consider a column of the feature matrix

$$\mathbf{H}^* \in \mathbb{R}^{N \times d}$$

TL;DR:

- A small modification on GCNII allows for **arbitrary** polynomial filter expressiveness.
- Underlying is a correspondence between **evaluating method** and **convolution stack**.

A Warm-up

$$\text{GCNII: } \mathbf{H}^{(\ell+1)} = \sigma(((1 - \alpha)\tilde{\mathbf{P}}\mathbf{H}^{(\ell)} + \alpha\mathbf{H}^*)((1 - \theta_\ell)\mathbf{W}^{(\ell)} + \theta_\ell \mathbf{I}))$$

$$\mathbf{H}^{(\ell+1)} = (1 - \alpha)\tilde{\mathbf{P}}\mathbf{H}^{(\ell)} + \alpha\mathbf{H}^*$$

Unfold it!

$$\mathbf{H}^{(K)} = \sum_{\ell=0}^K \hat{\alpha}_\ell \tilde{\mathbf{P}}^\ell \mathbf{H}^*$$

$$\hat{\alpha}_\ell = \begin{cases} \alpha(1 - \alpha)^\ell, & \ell < K \\ (1 - \alpha)^K, & \ell = K \end{cases}$$

Free the coefficients!

$$\mathbf{H}^{(\ell+1)} = \tilde{\mathbf{P}}\mathbf{H}^{(\ell)} + \alpha_\ell \mathbf{H}^*$$

Unfold it!

$$\mathbf{H}^{(K)} = \sum_{\ell=0}^K \alpha_{K-\ell} \tilde{\mathbf{P}}^\ell \mathbf{H}^*$$

Unfolding Process (1) corresponds to **Horner's Method!** An iterative method for numerical evaluating.

Horner's Method: Evaluates the value of $p(x)$ at x_0 where $p(x) = a_0 + a_1 x + \dots + a_n x^n$.

$$\begin{aligned} b_n &:= a_n, \\ b_{n-1} &:= a_{n-1} + b_n x_0, \\ &\vdots \\ b_0 &:= a_0 + b_1 x_0, \\ \Rightarrow p(x_0) &:= b_0. \end{aligned}$$

$$\begin{aligned} \mathbf{H}^{(0)} &= \alpha_0 \mathbf{H}^*, \\ \mathbf{H}^{(1)} &= \tilde{\mathbf{P}}(\alpha_0 \mathbf{H}^*) + \alpha_1 \mathbf{H}^*, \\ \mathbf{H}^{(2)} &= \tilde{\mathbf{P}}(\tilde{\mathbf{P}}(\alpha_0 \mathbf{H}^*) + \alpha_1 \mathbf{H}^*) + \alpha_2 \mathbf{H}^*, \\ &\vdots \\ \mathbf{H}^{(K)} &= \tilde{\mathbf{P}}(\dots(\tilde{\mathbf{P}}(\tilde{\mathbf{P}}(\alpha_0 \mathbf{H}^*) + \alpha_1 \mathbf{H}^*) + \alpha_2 \mathbf{H}^*) \dots) + \alpha_K \mathbf{H}^* \\ \Rightarrow \mathbf{H}^{(K)} &= \alpha_K \mathbf{H}^* + \alpha_{K-1} \tilde{\mathbf{P}}\mathbf{H}^* + \dots + \alpha_0 \tilde{\mathbf{P}}^K \mathbf{H}^* \\ &= \sum_{\ell=0}^K \alpha_{K-\ell} \tilde{\mathbf{P}}^\ell \mathbf{H}^*. \end{aligned}$$

Unfolding Process

(a) Comparison with other models equipped with different kinds of residual connections. Mean classification accuracies (\pm standard deviations) of twenty random splits are displayed. Besides the ClenshawGCN, all the results are taken directly from Luan et al. [28] and Lim et al. [26].

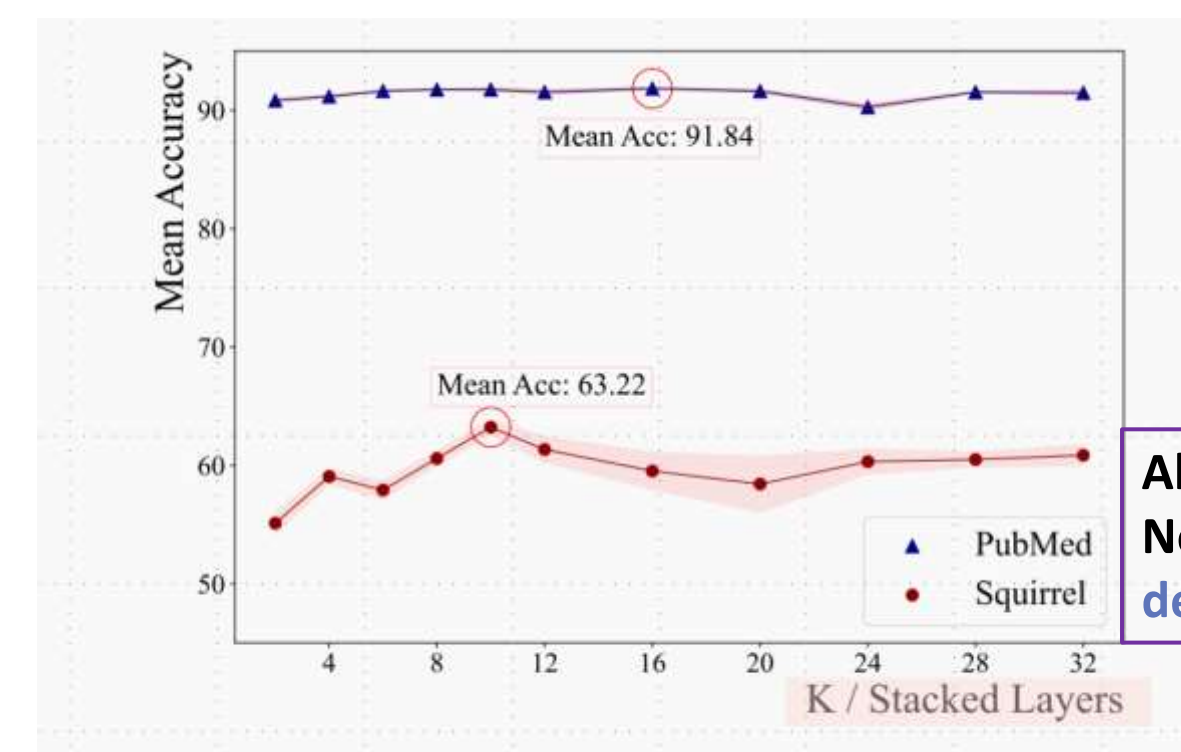
| Datasets | Chameleon | Squirrel | Actor | Texas | Cornell | Cora | Citeseer | PubMed |
|--------------------|-------------|------------|------------|------------|-------------|-------------|-------------|------------|
| [V] | 2,277 | 5,201 | 7,600 | 183 | 183 | 2,708 | 3,327 | 19,717 |
| MPL | 46.59±1.84 | 31.01±1.18 | 40.18±0.35 | 86.81±2.24 | 84.15±3.05 | 76.89±0.97 | 76.52±0.89 | 86.14±0.25 |
| GCN | 60.81±2.95 | 45.87±0.88 | 33.26±1.15 | 76.97±3.97 | 65.78±4.16 | 37.18±1.12 | 79.85±0.78 | 86.79±0.31 |
| GCNII | 63.44±0.85 | 41.96±1.02 | 36.89±0.95 | 80.46±5.91 | 84.26±2.13 | 88.46±0.82 | 79.97±0.65 | 89.94±0.31 |
| H ₂ GCN | 52.39±0.48 | 30.39±1.22 | 38.85±1.17 | 85.90±3.53 | 86.25±4.71 | 87.52±0.61 | 79.97±0.69 | 87.78±0.28 |
| MixHop | 36.28±10.22 | 24.55±2.60 | 33.13±2.40 | 76.39±7.66 | 60.33±28.53 | 65.65±11.31 | 49.52±13.35 | 87.04±4.10 |
| GCN+JK | 64.68±2.85 | 53.40±1.90 | 32.72±2.62 | 80.66±1.91 | 66.56±13.82 | 86.90±1.51 | 73.77±1.85 | 90.09±0.68 |
| ClenshawGCN | 69.44±2.06 | 62.14±1.65 | 42.08±1.99 | 93.36±2.35 | 92.46±3.72 | 88.90±1.26 | 80.34±1.26 | 91.99±0.41 |

v.s. Representative residual-enhanced spatial GNNs.

Large margin on hete-graphs

v.s. SoTA spectral GNNs

Competitive!

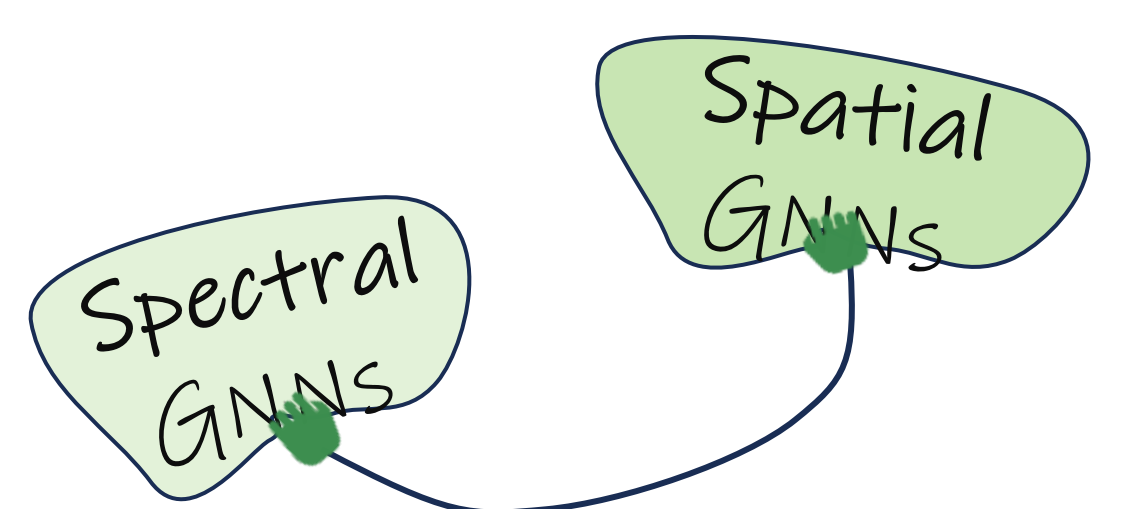


Experiments Node classification & Ablations

Ablation: No model degradation

| Ablation: Effectiveness of intertwining nonlinear transformations into polynomial filters. | Models | ClenshawGCN | ClenshawGCN(-Act) | Clenshaw(-Act-W) |
|--|-----------|--------------|-------------------|------------------|
| | Squirrel | 62.14 ± 1.65 | 61.55 ± 1.42 | 56.92 ± 2.13 |
| | Chameleon | 69.45 ± 2.12 | 67.29 ± 2.35 | 70.08 ± 2.43 |
| | PubMed | 91.99 ± 0.41 | 91.56 ± 0.46 | 91.27 ± 0.53 |
| | Penn94 | 85.38 ± 0.25 | 84.68 ± 0.56 | 84.39 ± 0.26 |

Take-Home Messages



- **Spatial GNNs** and **spectral GNNs** adopt different perspectives in utilizing graphs.

- **Spectral GNNs** are dominated by **polynomial filters** so far. SoTA polynomial filters can approximate **arbitrary polynomial functions** using **polynomial basis**.
- **Spatial GNNs** benefit from entangled non-linear transformations.

- We use **simple residual connections** to rewire the information flow, injecting **spectral characteristics** into a **message passing (spatial) backbone**, keeping the entangled transformations.

- The **stack of convolution blocks** aligns with the **iteration of Clenshaw's algorithm**.
- There is a special **negative residual**. The role of it is to use Chebyshev Basis.