Deep Networks as *hidden* Metric Learners

- N training instances: $x_1, \ldots, x_n, \ldots, x_N$

- Ground truth training labels: $y_1, \ldots, y_n, \ldots, y_N$

- Seek a function, $f : \mathbb{X} \to \mathbb{Y}$, to predict $\hat{y}_{N+1}$ for a new, unseen instance x_{N+1}, with minimal *distance* between $\hat{y}_{N+1}$ and y_{N+1}

- New view: Back-out a metric learner from the parametric deep network:
$f = c \circ g$, where $g : \mathbb{X} \to \mathbb{R}^M$, $c : \mathbb{R}^M \to \mathbb{Y}$, and $r \in \mathbb{R}^M$ is a dense representation of the input under the parametric model

- Sense in which: $f\left(x_{N+1}\right) \approx \beta + \sum_{n=1}^{N} \left(\tanh(f(x_n)) + \gamma \cdot y_n\right) \cdot w\left(||r_n - r_{N+1}||_2\right)$

I.e., a test prediction is approx. a distance-weighting (between "***exemplar*** representations) over the training set (model predictions & associated labels)

$w(\cdot)$ is a function of the distance between representations
(Relatable to instance-based learning, kernel methods, …)

- Enables interpretable/introspectable decision rules & various analyses (hence, "***auditing***"): E.g., only admit true positive (TP) matches:
$\hat{y}_{N+1} = f(x_{N+1}) \cdot \left[f(x_{N+1}) = f(x_n) \wedge f(x_n) = y_n\right] + NULL \cdot \left[f(x_{N+1}) \neq f(x_n) \vee f(x_n) \neq y_n\right]$, where $n = \underset{n \in \{1,\ldots,N\}}{\arg\min}\ ||r_n - r_{N+1}||_2$

- Enables updatability/adaptability:

  - Label changes: $y'_n = y_n + \Delta_n$

  - Data additions (a.k.a., continual/lifelong learning):
  $\mathbb{D}^N = \left\{(x_1, y_1), \ldots, (x_N, y_N)\right\}$ becomes $\mathbb{D}^{N'} = \left\{(x_1, y_1), \ldots, (x_N, y_N), \ldots, (x_{N'}, y_{N'})\right\}$

  - New lightweight models over representations (e.g., using data additions): $c' : \mathbb{R}^M \to \mathbb{Y}'$

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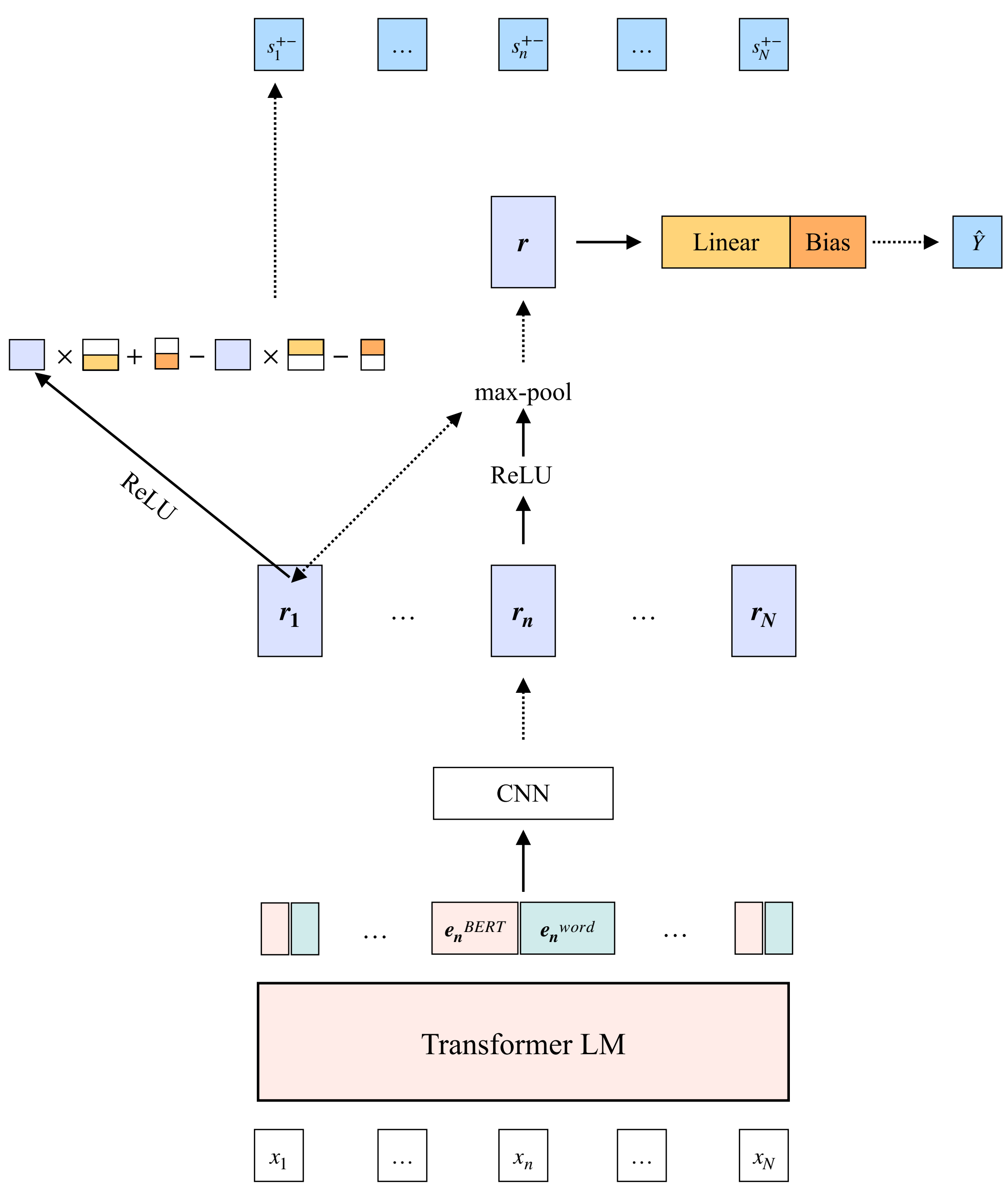
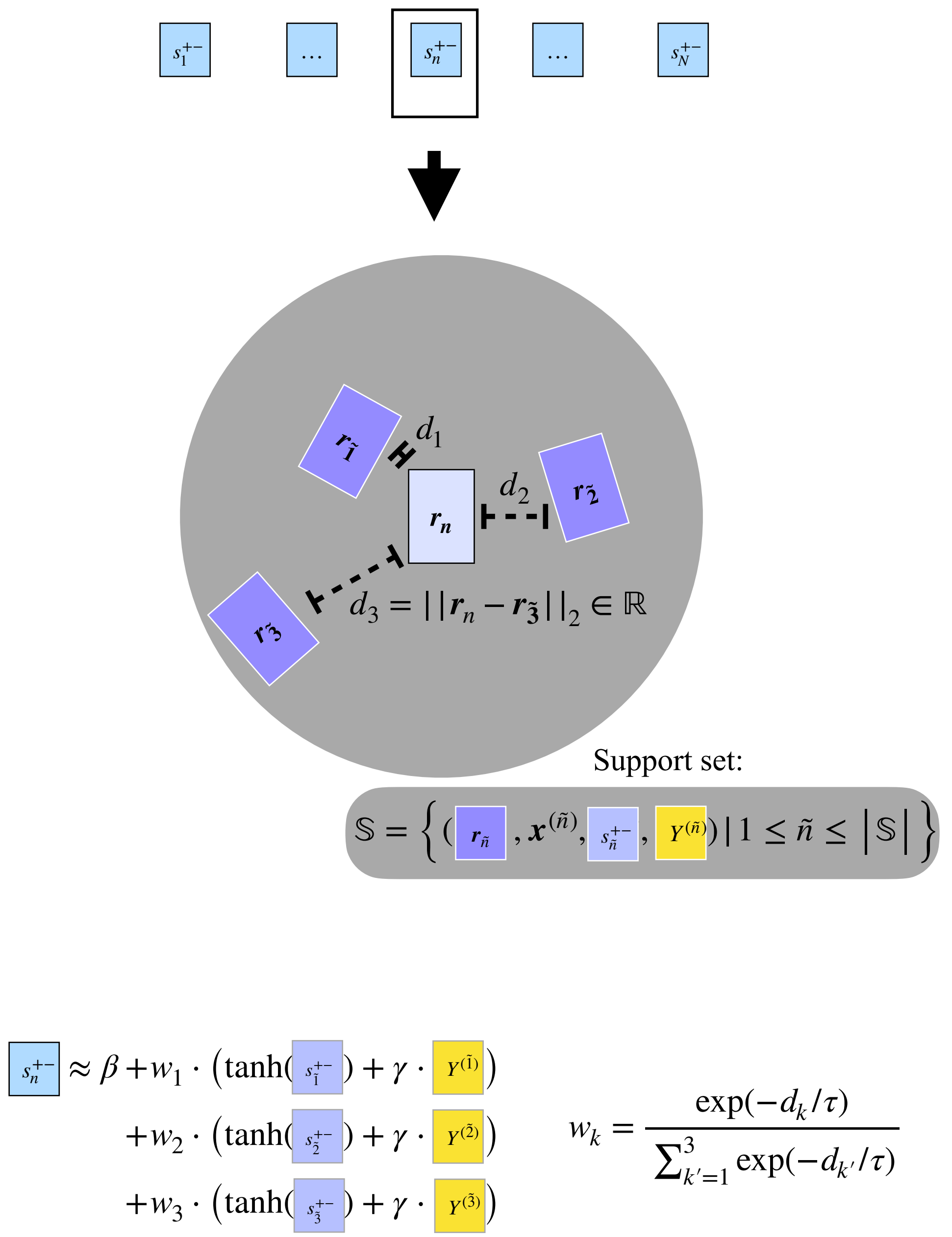
Horizontal (across the input) & *Vertical* (across the support set) Model Decompositions

Sequence Labeling via a Convolutional Decomposition



K-NN Approximation



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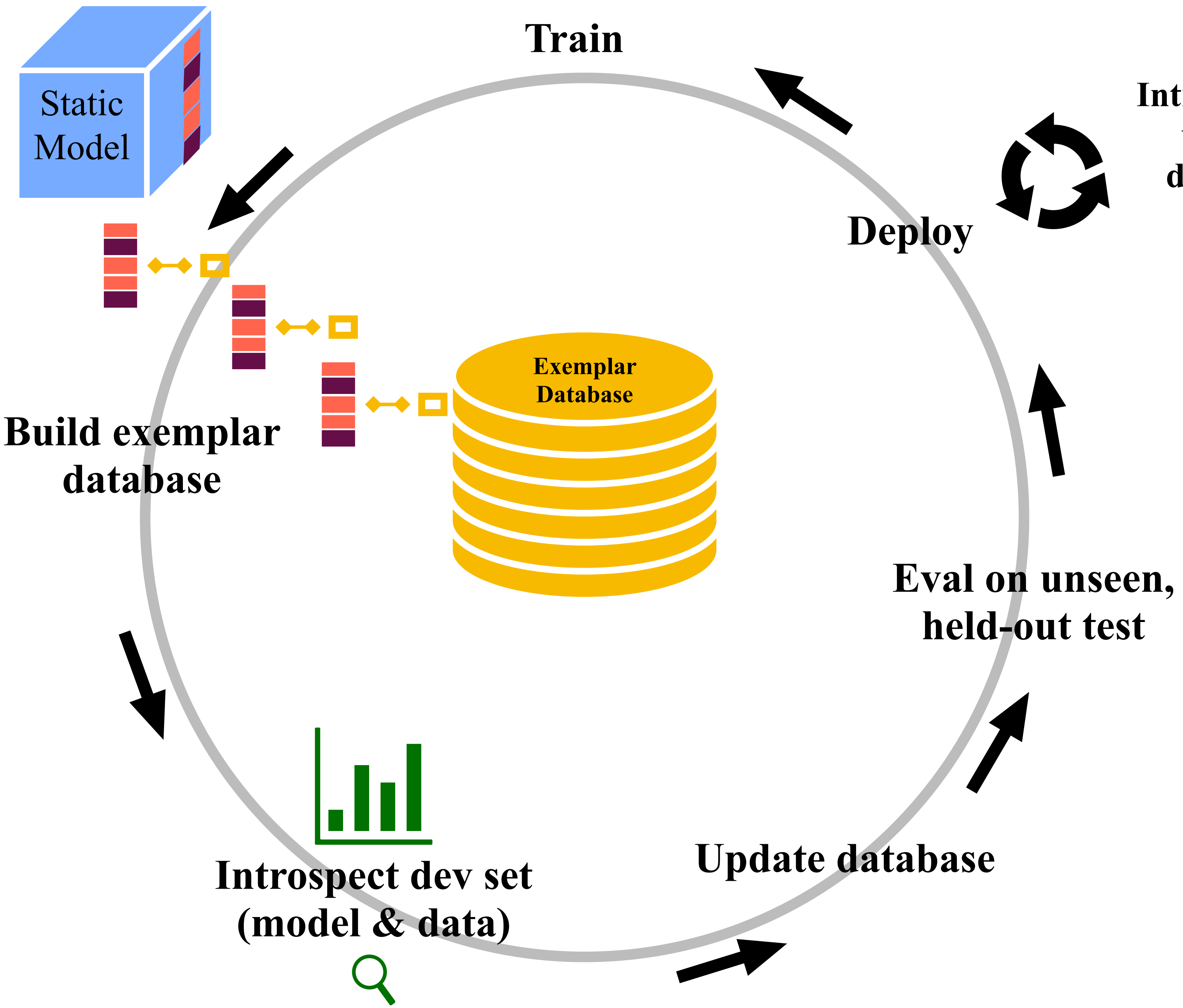
Exemplar Auditing Lifecycle



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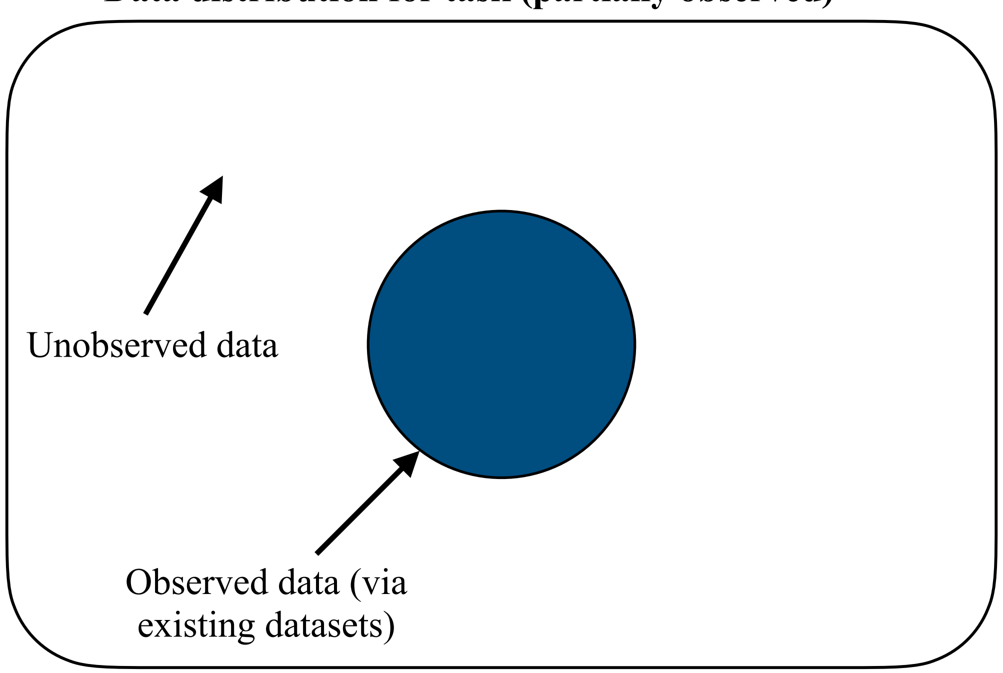
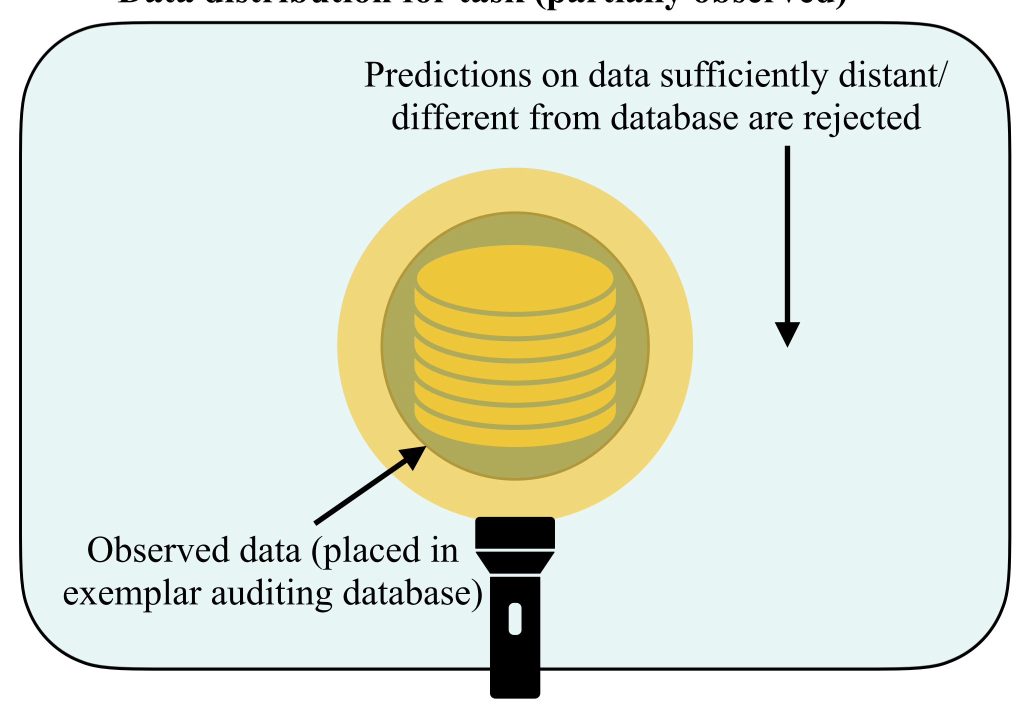
Out-of-Domain Settings

- Pre-train with as much data as possible
- Add as much data as possible to the database, including data not seen in training
  - Corral the in-domain space, around the ball of the observed data
  - Never predict over out-of-domain data in high-risk settings. Instead: Rearrange the deployment to handle non-admitted predictions.

Data distribution for task (partially observed)



Data distribution for task (partially observed)



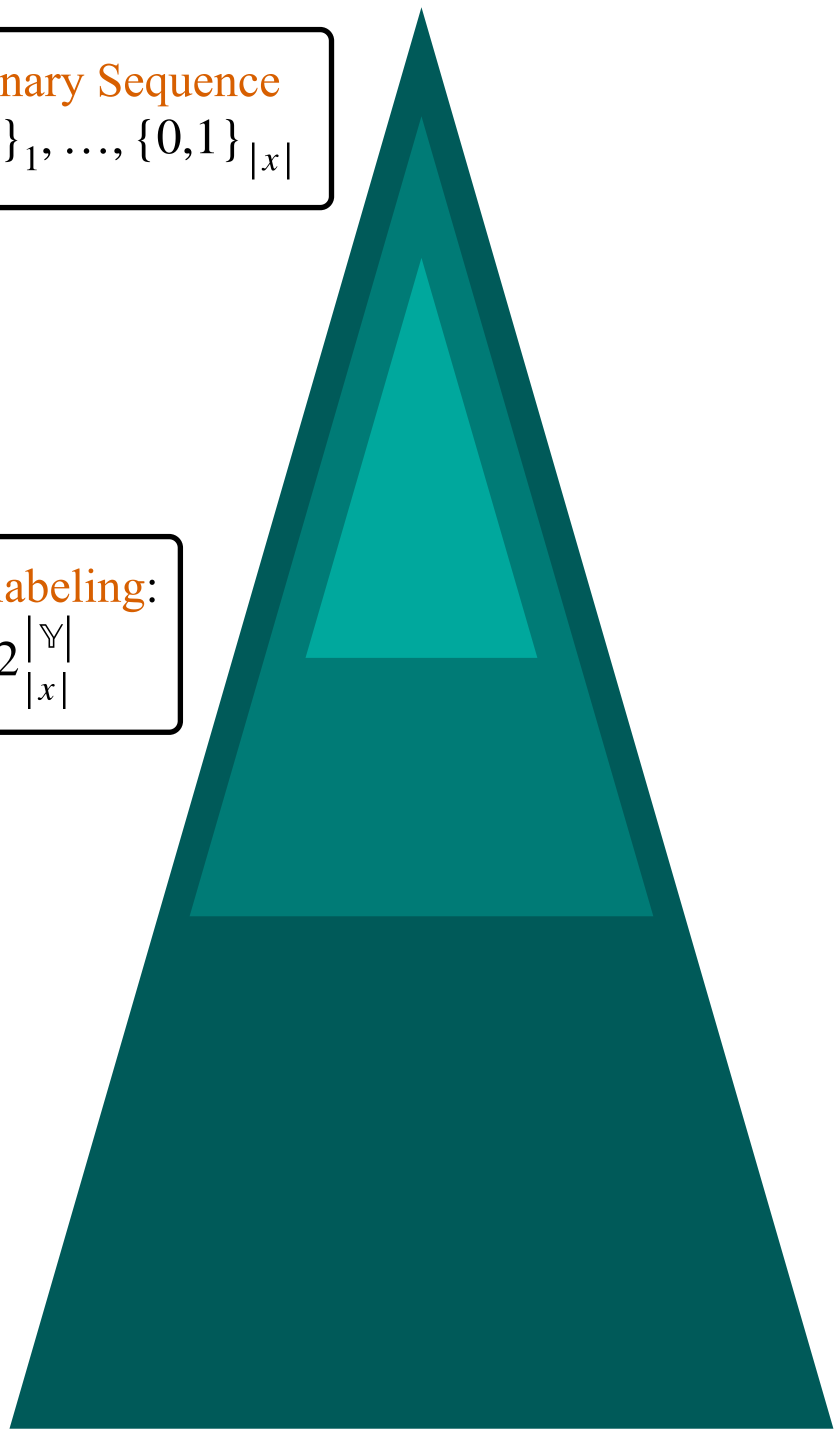
Implementations

Unique side effect: Binary Sequence labeling: $f: \mathbb{X} \rightarrow \{0,1\}_1, \ldots, \{0,1\}_{|x|}$

- Binary classification: $f: \mathbb{X} \rightarrow \{0,1\}$
  - "Detecting Local Insights from Global Labels: Supervised & Zero-Shot Sequence Labeling via a Convolutional Decomposition"

Multi-label sequence labeling: $f: \mathbb{X} \rightarrow 2_1^{|\mathbb{Y}|}, \ldots, 2_{|x|}^{|\mathbb{Y}|}$

- Multi-label classification: $f: \mathbb{X} \rightarrow 2^{|\mathbb{Y}|}$
  - "Exemplar Auditing for Multi-Label Biomedical Text Classification"
- Retrieval-classification: $f: \mathbb{X} \times \mathscr{D} \rightarrow \left\langle \{0,1,2\},\ 2^{|\mathbb{D}|} \right\rangle$
  - "Coarse-to-Fine Memory Matching for Joint Retrieval and Classification"



Memory Matching Search

- Approach (*high-level*): Run the same shared network, g, over all of Wikipedia, $\mathbb{D}$, caching the representations, & then perform search by matching the query representation with progressively built-up support sequences

q = `Query sequence`

s = `Support sequence`

A Wikipedia sentence

Search Level 1 $g(q) = r_q \in \mathbb{R}^M \longleftrightarrow g(s_1) = r_{s_1} \in \mathbb{R}^M$

$\vdots$

$g\left(s_{|\mathbb{D}|}\right) = r_{s_{|\mathbb{D}|}} \in \mathbb{R}^M$

$r_{s_1}, \ldots, r_{s_{|\mathbb{D}|}}$ can be cached

$s_i \in \mathbb{D}$

Set of K nearest Wikipedia sentences

$$s'_k \in \arg K \min_{s_i} ||r_q - r_{s_i}||_2$$

Search Level 2 $r_q \longleftrightarrow g\left((q, s'_1)\right) = r_{(q,s'_1)} \in \mathbb{R}^M$

$\vdots$

$g\left((q, s'_K)\right) = r_{(q,s'_K)} \in \mathbb{R}^M$

Set of Z nearest Wikipedia sentences from Search Level 2

$$s''_z \in \arg Z \min_{s'_k} ||r_q - r_{(q,s'_k)}||_2$$

Search Level 3

$$\hat{y} = \arg\min_{y \in \{\text{Supports, Refutes, Unverifiable}\}} ||r_q - r_{(y,q,s''_1,\ldots,s''_Z)}||_2$$

$\hat{y}$ is the label prediction

$\{s''_1, \ldots, s''_Z\}$ is the set of Wikipedia support sentences

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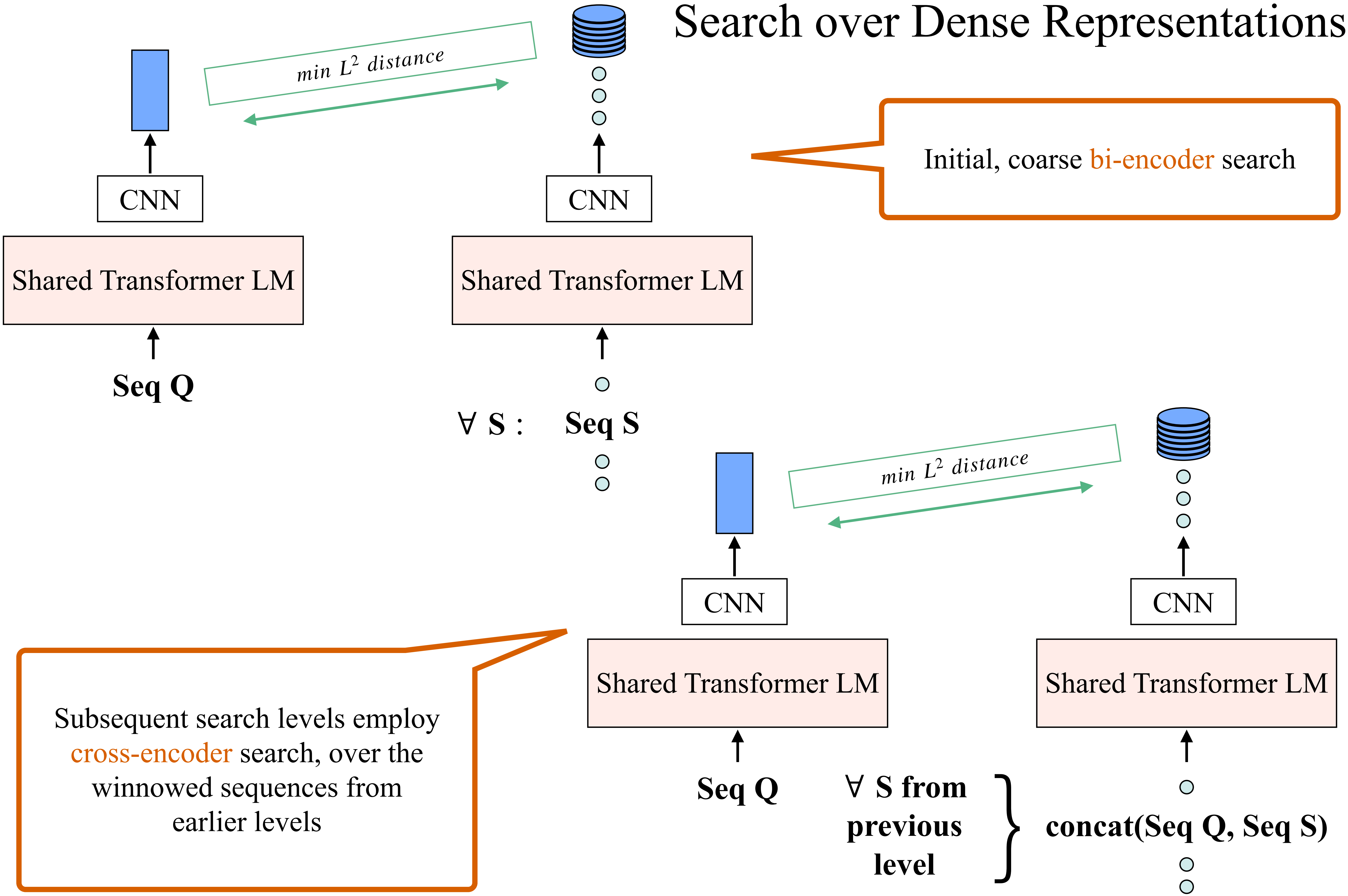
An End-to-End Retrieval-Classification Model via a Coarse-to-Fine Search over Dense Representations



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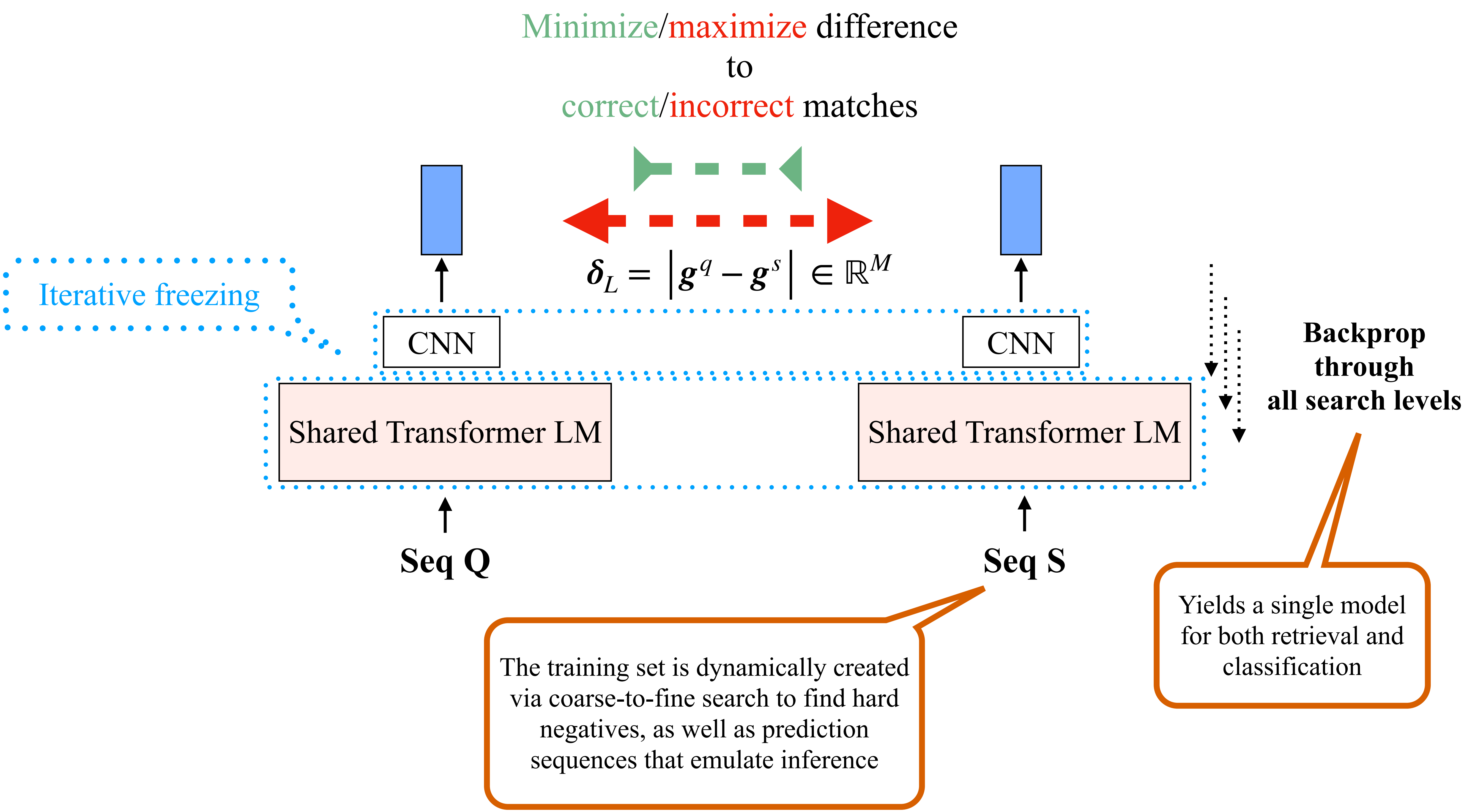
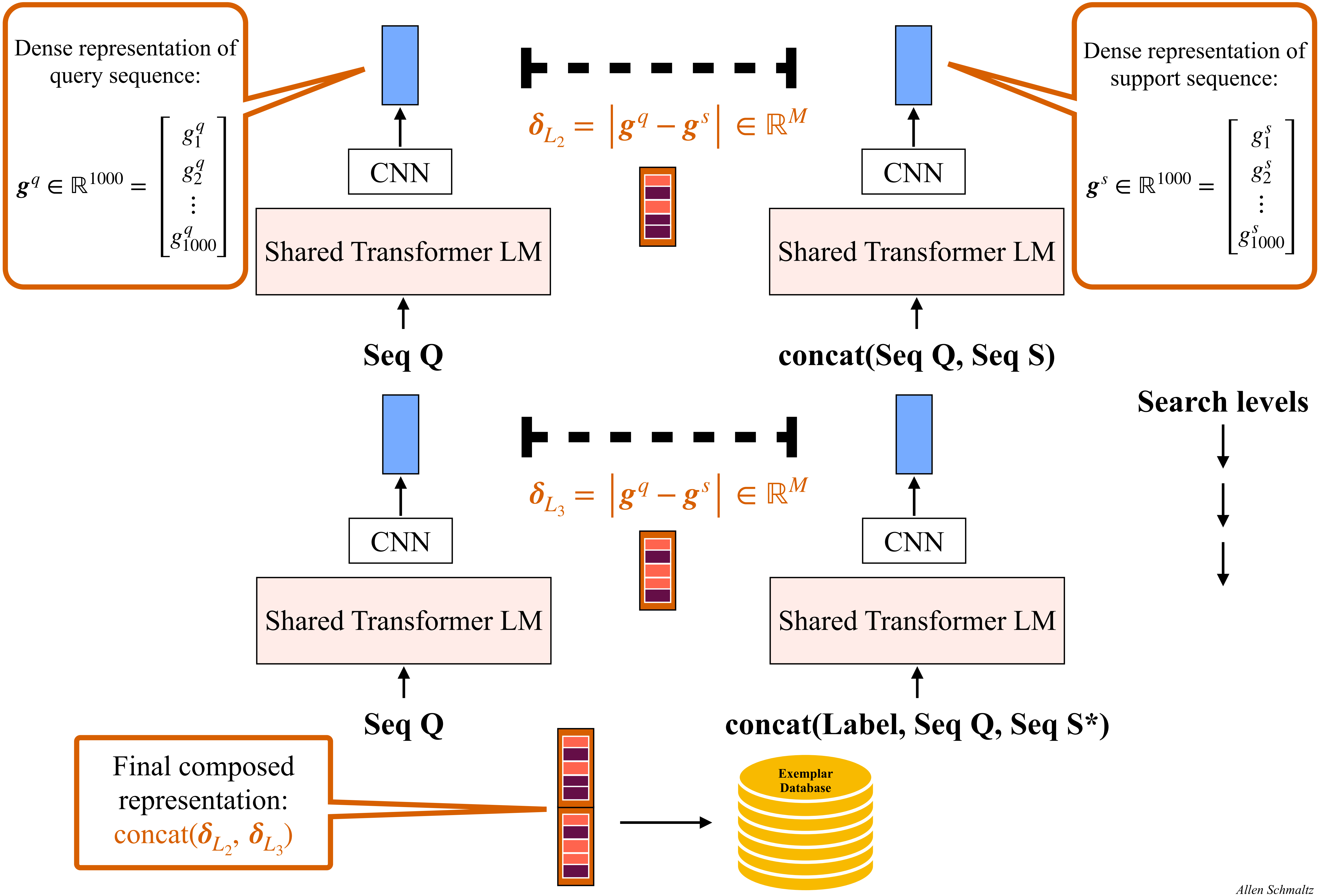
Joint Retrieval and Classification Training

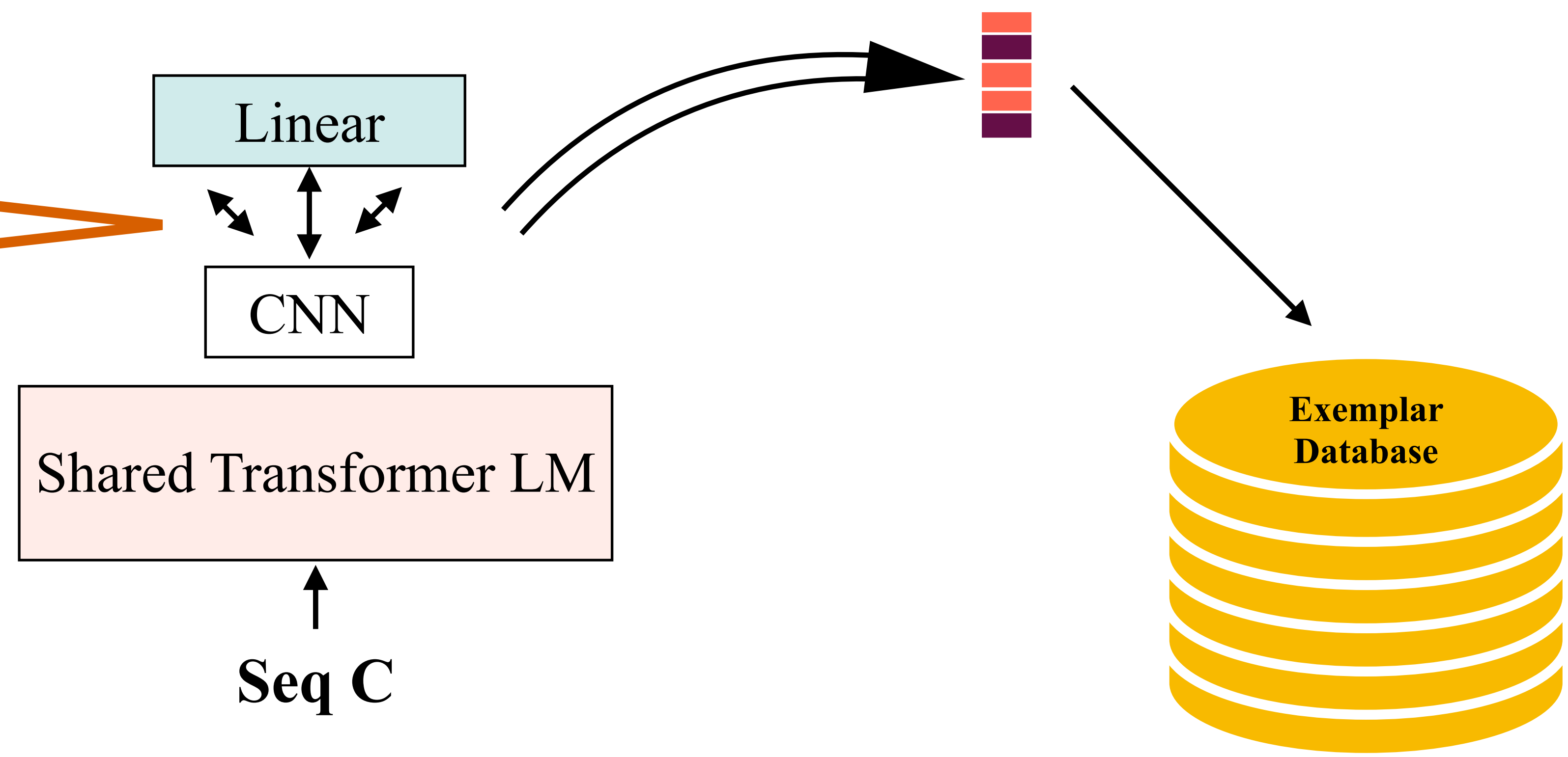


Multi-Sequence Representation Composition for Exemplar Auditing

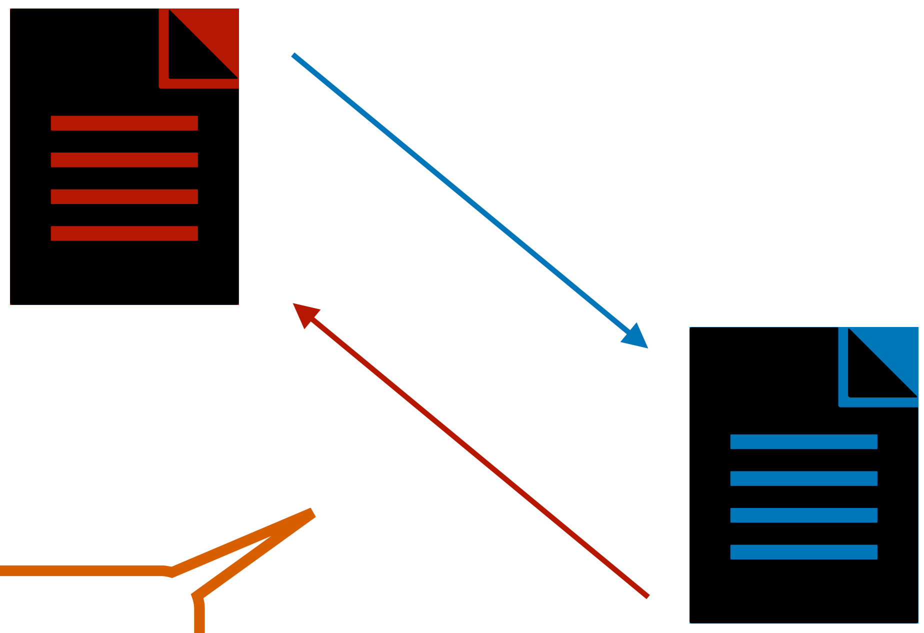


Token-Level Representations for Exemplar Auditing

Identify the dense representation of a token-level feature using [multi-] binary labeling via a convolutional decomposition (optionally, with priors to encourage/discourage particular features)

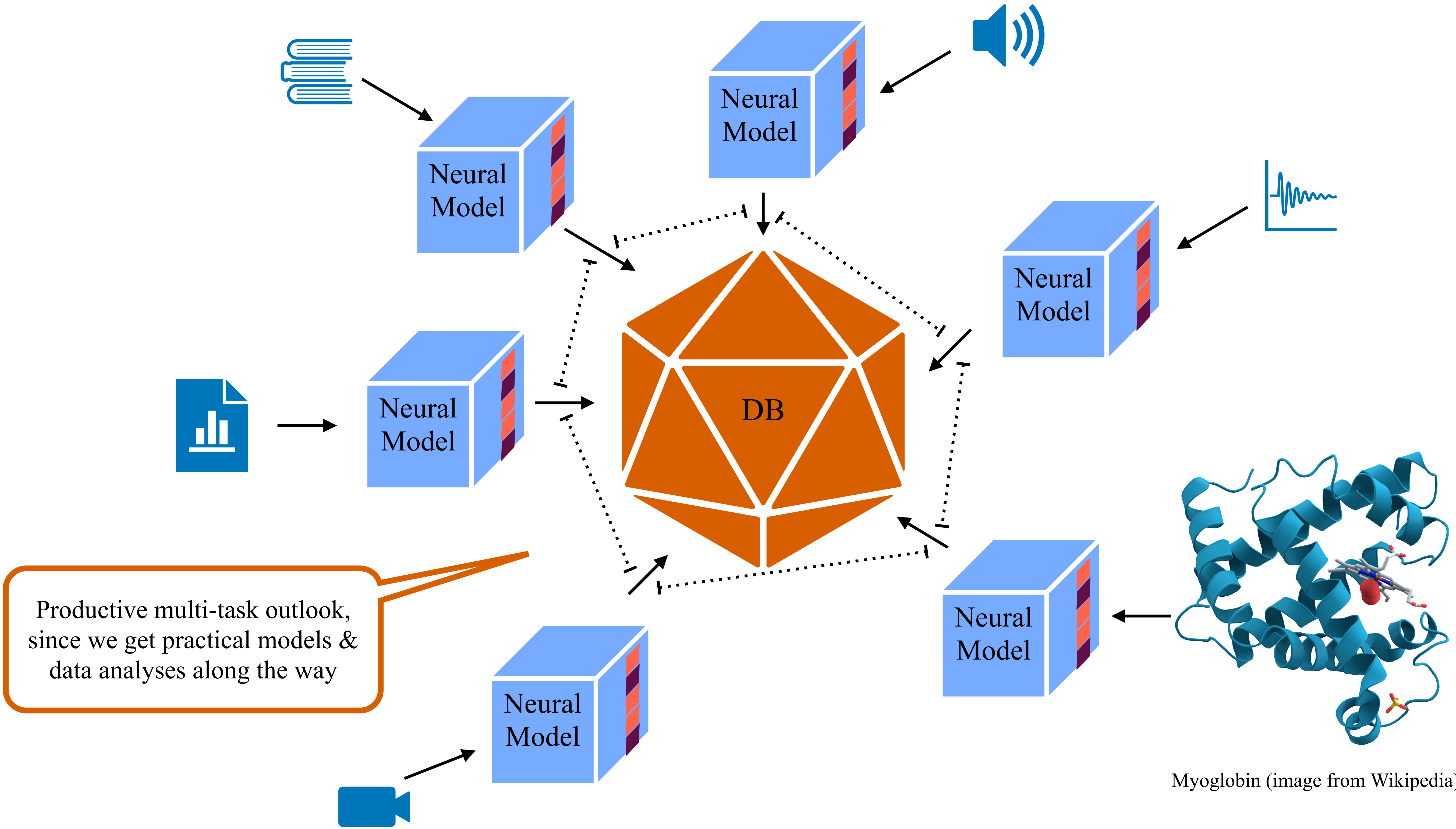


Extractive, Comparative (Feature-wise) Summarization



With facility over features, relating a global prediction to individual sequence elements, we can readily score, examine, & compare salient subsequences across correct & incorrect predictions for each class

Prospective Outlook: Interlocking distance constraints across input modalities and tasks via a single, shared model and a dense database…



Myoglobin (image from Wikipedia)

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