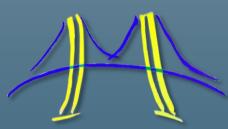
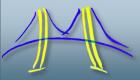
PRODUCTIVE GMM TRAINING WITH SEJITS FOR SPEAKER DIARIZATION

Katya Gonina, Henry Cook, Shoiab Kamil, Gerald Friedland, Armando Fox, David Patterson

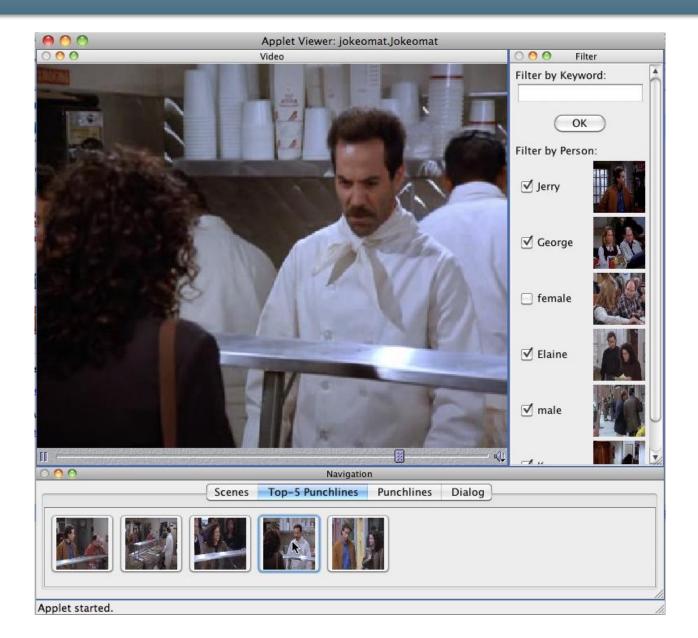
ParLab Retreat, June 2, 2011



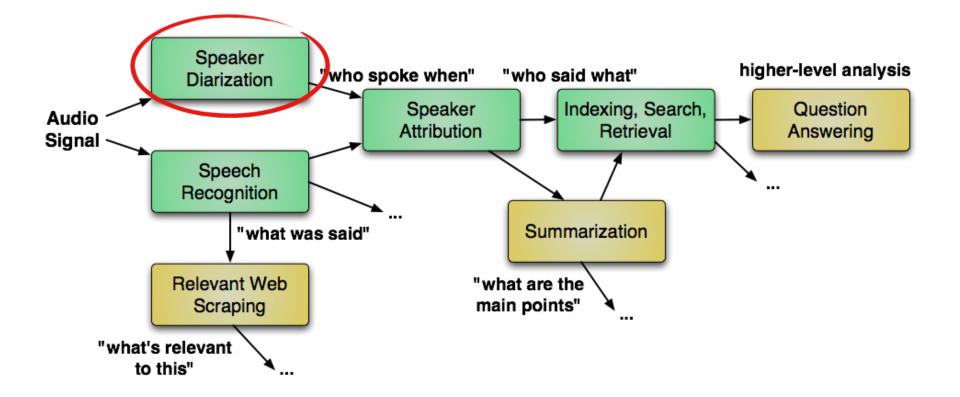


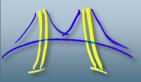


The Meeting Diarist



Components of the Meeting Diarist





Speaker Diarization

Audio track:

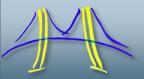
Segmentation:

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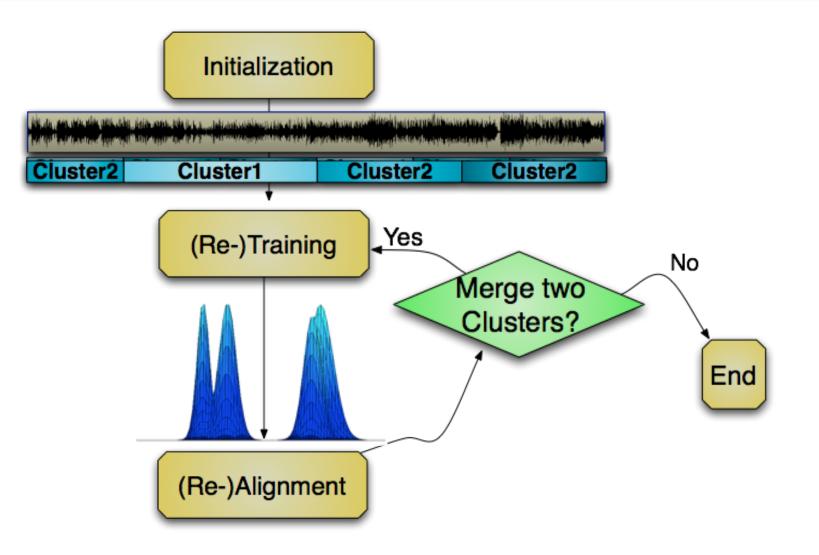
Clustering:

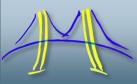
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Estimate "who spoke when" with no prior knowledge of speakers, #of speakers, words, or language spoken.



Speaker Diarization: Core Algorithm

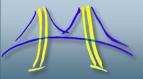




- Five versions (so far):
- Initial code (2006): 0.333 x realtime (i.e., 1h audio = 3h processing)
- Serially optimized (2008): 1.5 x realtime
- Parlab retreat summer 2010: Multicore+GPU parallelization: 14.3 x realtime
- Parlab retreat winter 2011: GPU-only parallelization 250 x realtime (i.e., 1h audio = 14.4sec processing)
 - -> Offline = online!

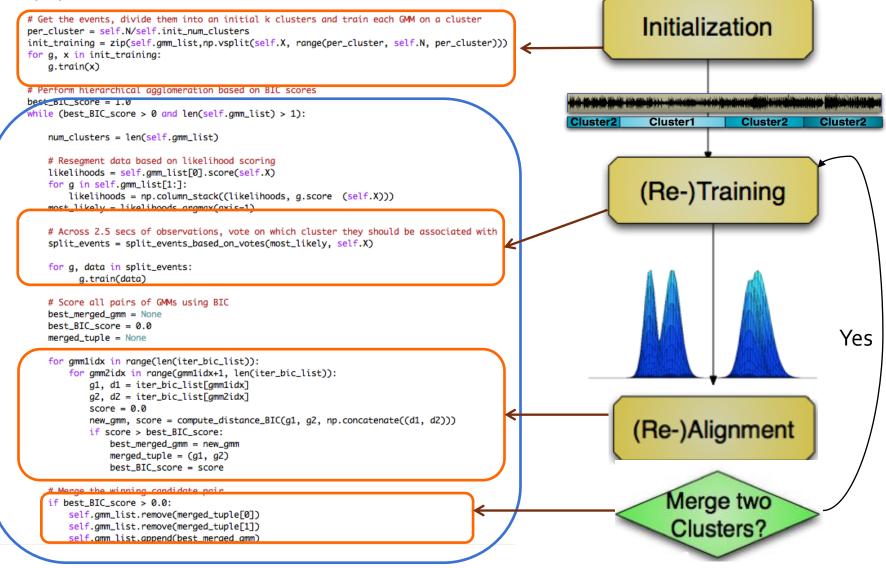
Parlab retreat summer 2011: SEJITized! [1]

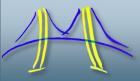
[1] H. Cook, E. Gonina, S. Kamil, G. Friedland, D. Patterson, A. Fox. CUDA-level Performance with Pythonlevel Productivity for Gaussian Mixture Model Applications. USENIX HotPar Workshop, 2011.



Speaker Diarization in Python

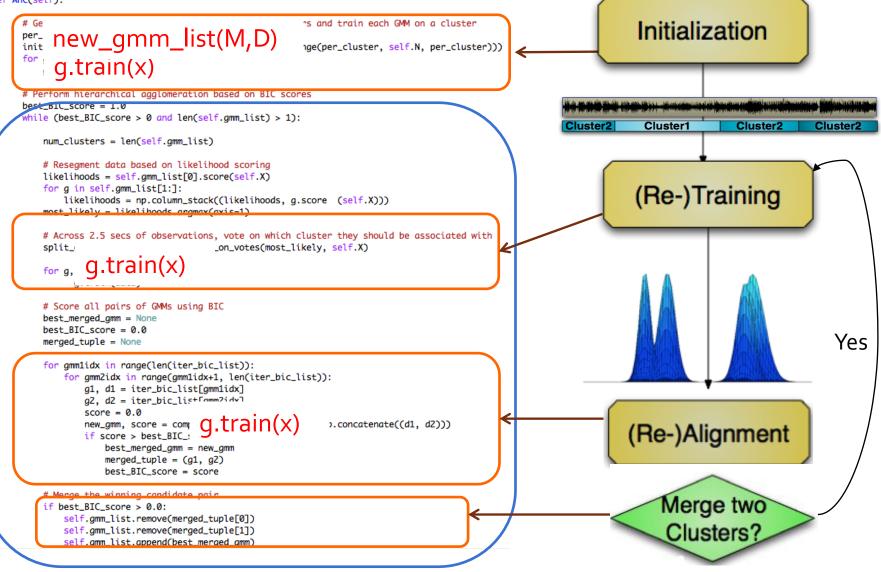
def AHC(self):

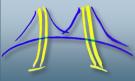




Speaker Diarization in Python



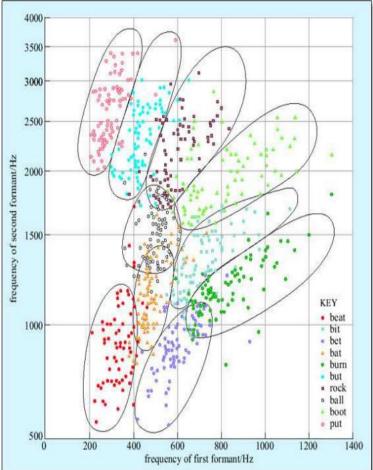




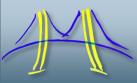
Gaussian Mixture Models & Training



- GMM probabilistic model for clustering (audio) data
 - Assumes the distribution of observations follows a set (mixture) of multidimensional Gaussian distributions
 - Each Gaussian in the mixture has a mean (*n*) and a covariance (*s*) parameters
 - Gaussians in the mixture are weighted with weight *D*



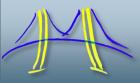
Example GMM in two dimensions (Source: Dan Klein, UCB)



GMM Training using EM Algorithm

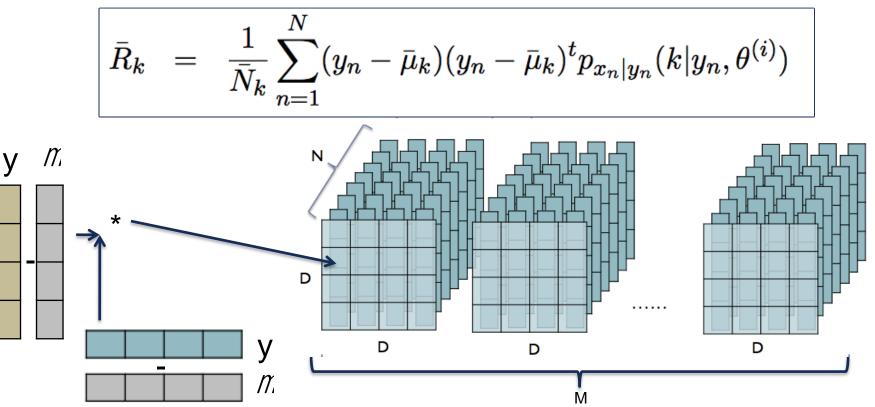
- Given a set of observations/events find the maximum likelihood estimates of the set of Gaussian Mixture parameters (*n*, *S*, *D*) and classify observations
- Expectation Maximization (EM) Algorithm
 - E step
 - Compute probabilities of events given model parameters
 - M step
 - Compute model parameters given probabilities
 - weights, mean, covariance matrix
 - Iterate until convergence

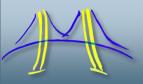
Covariance matrix – most computationally intensive step



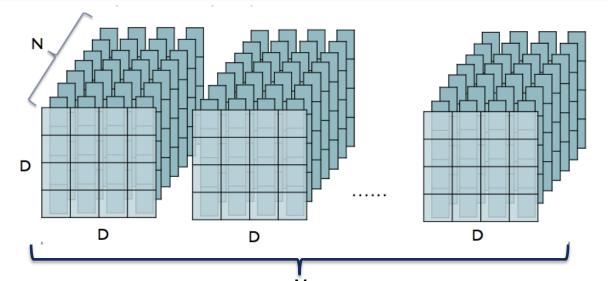
Covariance Matrix Computation

- N number of feature vectors, ~10K-100K
- D feature vector dimension (19 for speaker diarization), ~10-100
- M number of Gaussian components, ~1-128
- Matrix is symmetric only compute the lower DxD/2 cells

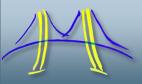




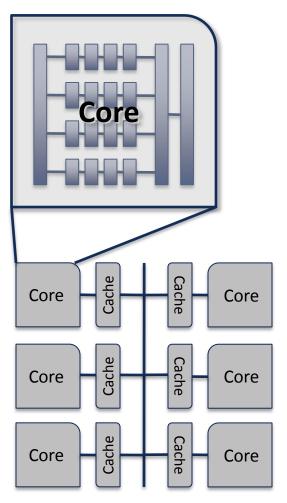
Covariance Matrix Computation



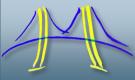
- Opportunities for parallelism (independent computations):
 - Each component's covariance matrix (M)
 - Each cell in a covariance matrix (DxD/2)
 - Each event's contribution to a cell in a covar matrix (N)
- -> Multiple code variants to perform the same computation in different ways



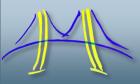
Manycore Parallel Platform



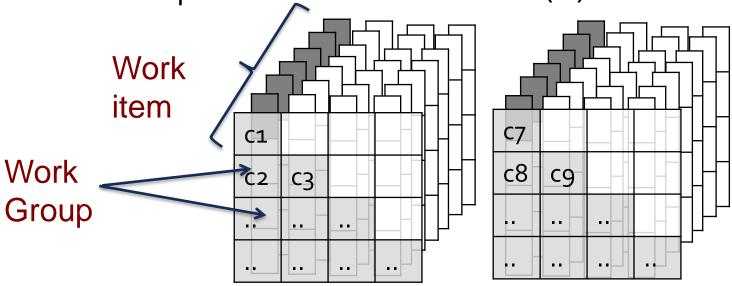
- Two levels of parallelism:
 - Work-groups parallelized across cores (CUDA threadBlock)
 - Work-groups' work-items executed on a single core, utilizing within-core parallelism (CUDA thread)
- Per-core local memory



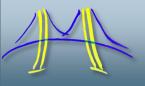
Code Variants



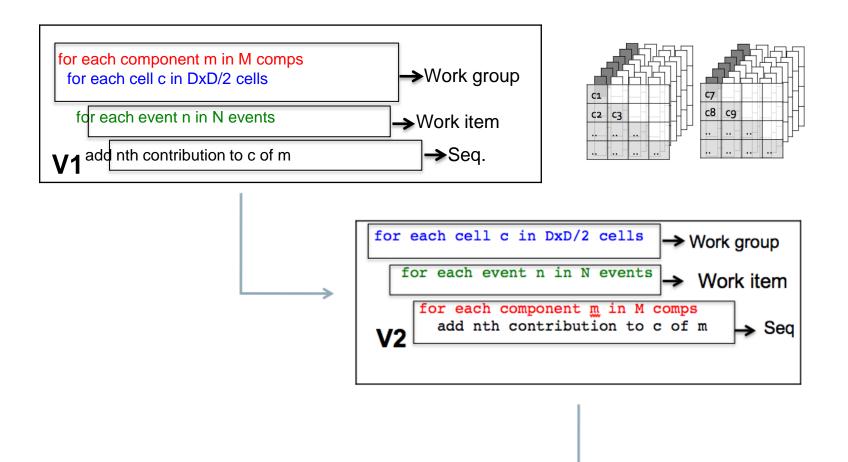
- Code variant 1:
 - 2D grid of work groups M x D x D/2
 - Each work group is responsible for computing one cell in the covariance matrix for one component
 - Work item parallelization over events (N)

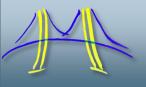


[2]. A. D. Pangborn. Scalable data clustering using gpus. Master's thesis, Rochester Institute of Technology, 2010.

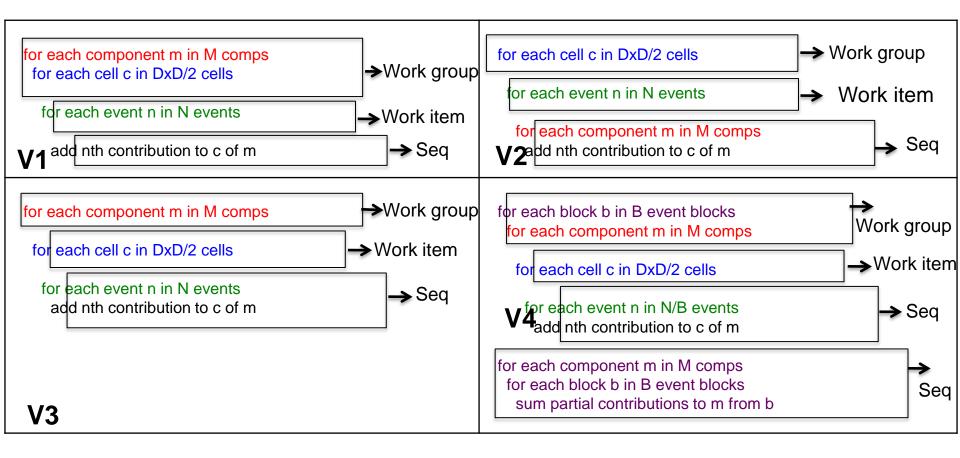


Covariance Matrix Computation – Code Variants

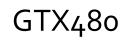


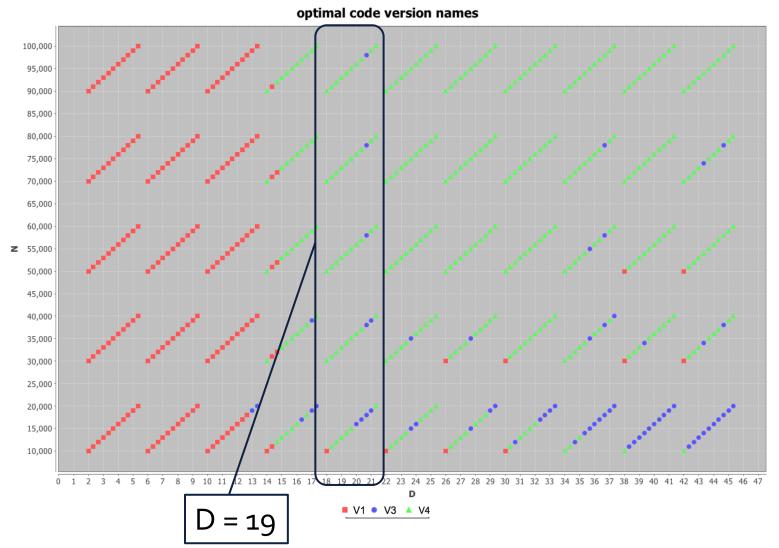


Covariance Matrix Computation – Code Variants Summary



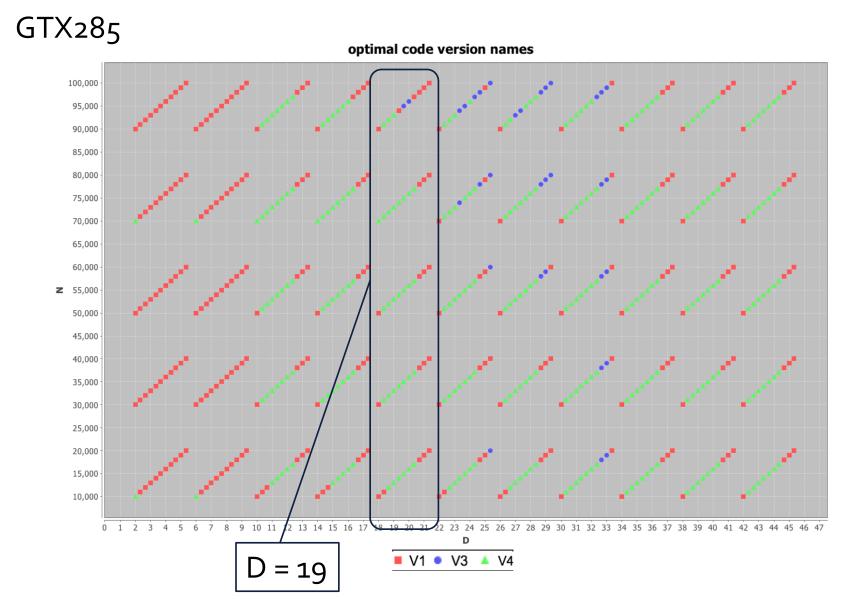
Results – Code Variant Performance



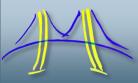


19/42

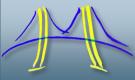
Results – Code Variant Performance



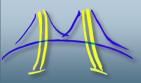
20/42



- Using best-performing code variant gave 32% average improvement in matrix computation time compared to always using original hand-coded variant (D: 1 to 36, M: 1 to 128, N: 10K to 150K)
- Performance gap increases with larger problem sizes (75.6% for D=36, M=128, N=500,000)

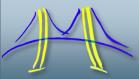


Specialization

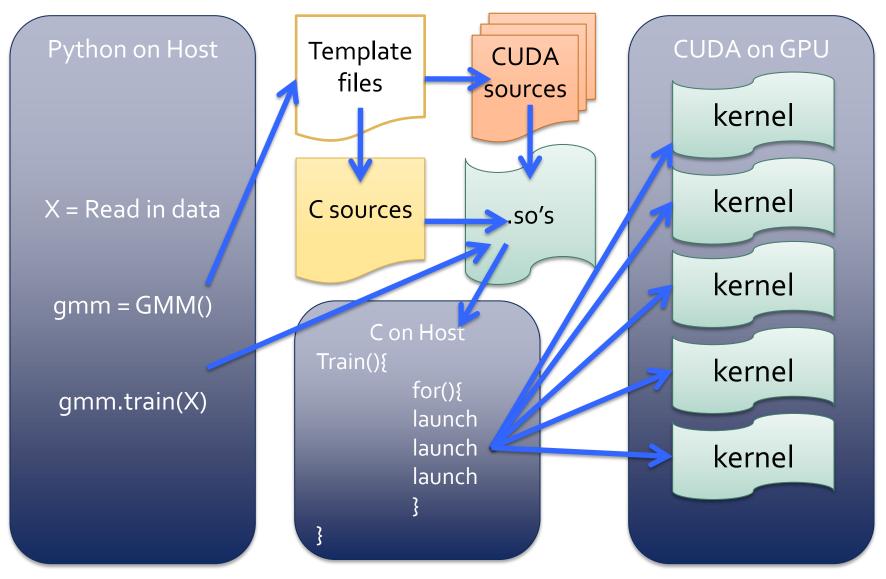


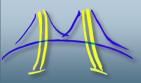
Specialization with ASP

- Given:
 - Problem Dimensions (N, D, M)
 - Platform Parameters (targeting Nvidia GPUs)
 - Core count, shared memory size, SIMD width...
- Automatically select:
 - Optimal code variant
 - Optimal parameters (block size, number of blocks) for that parallelization strategy



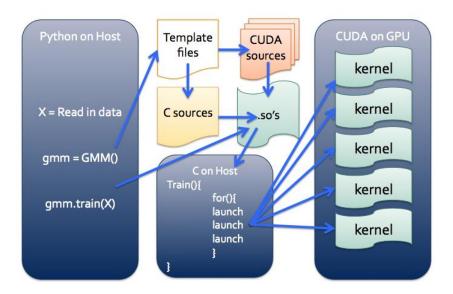
SEJITS Framework: Overview

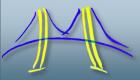




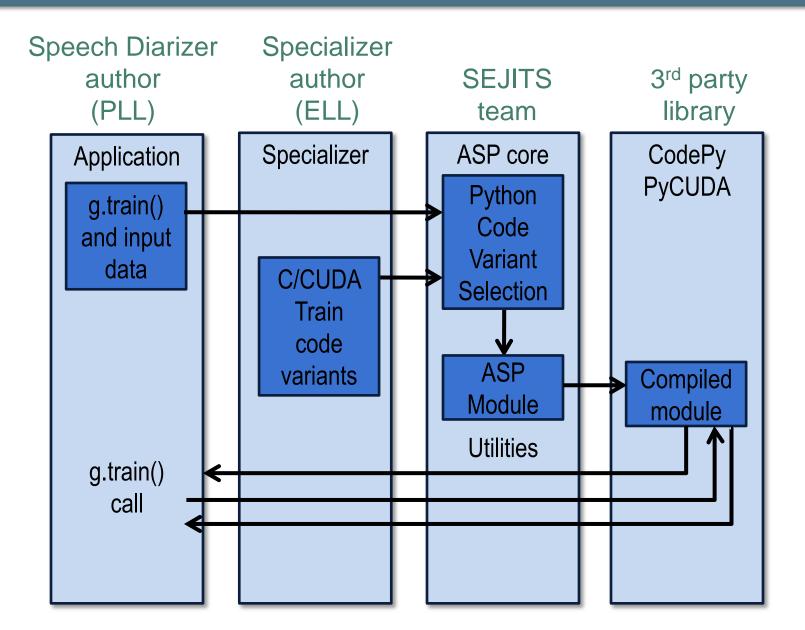
SEJITS Framework

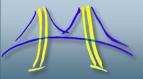
- Python code that handles application
 - Manipulates problem data, determines learning targets
- C/CUDA code that runs quickly
 - Allocates GPU memory
 - Performs main EM iterative loop
- Specializer (ASP)
 - Selects appropriate code variant (from history)
 - Pulls in the template for the code variant, parameterizes it and compiles to binary



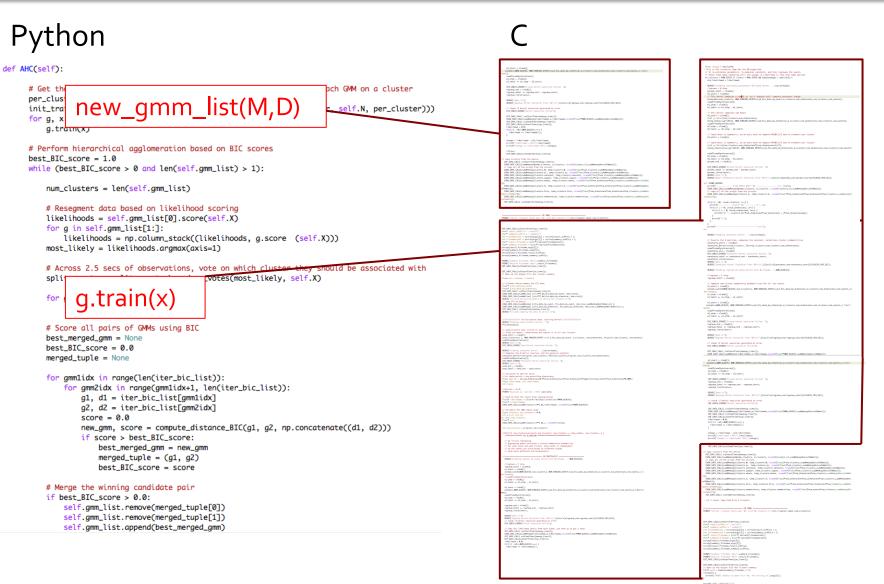


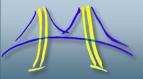
Separation of Concerns





Speaker Diarization in Python





Speaker Diarization in Python

Python

self.gmm_list.append(best_merged_gmm)

def AHC(self):

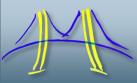
Get the events, divide them into an initial k clusters and train each GMM on a cluster XT_DEDC_EBB0C'E-step formel execution fell
vgraup_sold = Clock();
vgraup_sold = regraup_and = regraup_start; per_cluster = self.N/self.init_num_clusters init_training = zip(self.gmm_list,np.vsplit(self.X, range(per_cluster, self.N, per_cluster))) CRT_SAVE_CALL(excitantTimerQuee CRDA_SAVE_CALL(cuddMinepy(TikeTith CRT_SAVE_CALL(cudStarTimerQueep CRT_SAVE_CALL(cudStartTimerQuee) for g, x in init_training: g.train(x) idelifeed = 0.0; ending i=0.1delifeed(BLBCCS(1++) { likelifeed += likelifeed(2); chorge = likel printf("likeli # Perform hierarchical agglomeration based on BIC scores best BIC score = 1.0 while (best_BIC_score > 0 and len(self.gmm_list) > 1): num_clusters = len(self.gmm_list) # Resegment data based on likelihood scoring 15x LOC likelihoods = self.gmm_list[0].score(self.X) for g in self.gmm_list[1:]: likelihoods = np.column_stack((likelihoods, g.score most_likely = likelihoods.argmax(axis=1) reduction # Across 2.5 secs of observations, vote on which split_events = split_events_based_on_votes(most_likely for g, data in split_events: g.train(data) # Score all pairs of GMMs using BIC best_merged_gmm = None best_BIC_score = 0.0 merged_tuple = None for gmm1idx in range(len(iter_bic_list)): for gmmZidx in range(gmm1idx+1, len(iter_bic_list)): //epsilon = 1e-6; g1, d1 = iter_bic_list[gmm1idx] // Used to held the result free regrap kernel flact likelineast = (flact) mellection(f(flact)%40, flact d.likelineas; flact d.likelineas; flact d.likelineas; g2, d2 = iter_bic_list[gmm2idx] score = 0.0E (Marconstruction) Flash 4(4) CBA(MPE,DEL(combAlloc((cond*)) M(4), sizes((f)) new_gmm, score = compute_distance_BIC(g1, g2, np.concatenate((d1, d2))) if score > best_BIC_score: // de initial regresping
// Repropring mean solutions a cluster membership probability
// for each sent and add cluster. Each sumt is independent,
// is the events are distributed to different blocks
// ran bases different mitoprocessamy best_merged_gmm = new_gmm merged_tuple = (g1, g2)//regroup = E step regroup,stort = clack(); cl_start = clack(); esteplocation(NMM_BLOOK_P best_BIC_score = score # Merge the winning candidate pair s2_start = clash(); esteplocalM_RCCCS_NM_THEADS_CC if best_BIC_score > 0.0: self.gmm_list.remove(merged_tuple[0]) self.gmm_list.remove(merged_tuple[1]) BEBBC/Mare.V/% BEBBC/Sepress Reveal Discriming Disc: Ministry() // shock of Served second segment follow(); OK.DBC/SERBEC/Served second second follow();

That couple = sph(1ard); () This is the interviewing large for the DF algorithm. () If the entries promoting, the execution consistent, and then regression the exection () The entries these repeating and the sharing or illutional is large them same sphill with (form < 400, HTMS if (form < 400, HTMS M6.600(couple) > sphillon)) (shift)(form < 400, HTMS if (form < 400, HTMS M6.600(couple) > sphillon)) (Version of the second s estep.meret.up cudsfireadlynchrenize(); m2_viep = rlash(); m2_teta1 += m2_stop - m2_viort; // Covariance is symmetric, so we only need to compu-s3_start = clack(); // Environce is symmetric, so we only need to compute MP(N-1)/2 and gridDist(yww.sineters.yww.simemiares(yww.simemiares()/2)) mater.coverience.org/10/10, NM. IMEMIS. MIDboold. Frs. dots.bu. PRENE MATRIES _____ exact a stark(); constarts,kernel.ecoux.ctu subtreat/putrestar(); constarts,end = clack(); cdt, perco_tomer____ constarts_totel() == constarts, constarts_totel() == constarts, constarts_totel(); EBB(Clane.\n^2); EBB(Clane.\n^2); 00000Clineking regroup //regrave = E step regrave_stert = clock(); // Compute rem cluster membership probabilities for all the events d_start = class(); extenicod(d)DBC.BLOCG.tum.clusters).WBC.DBLASS.ESTEPso(d.frs.d) elitech); sl_step = cleck(); sl_tetal -- sl_step - sl_stert; e2.stort = clock(); estaple=NAM_RECES, NAM_THERES.FT MERS("done.in"); MERS("Reprose Rennel Disection Time: Bfurio",(// check if kernel execution generated on error CHT_DECK_ENDEC_Formal execution fulled"): s2_utars = clack(); synsploceMM_R.COS, MM_THEAD, CSTD>>>> OIT, OEOC, DRBR("T-stop Kinnel execution fails regress, and = stack[0] regress, teactions.end - regress, start; regress, teactions+: MERS("done.\n"); MERS("fograup Korvel Disrutian Time: MFv/of,(change = likelihood - ald_likelihood; prist("likelihood - NTV",likelihood); scient("likelihood - NTV",likelihood); (terms) NF WF (ALL(outStepTimer(coultimer)) (3)) OSMA_SWE_CALL(suddhimopy(clusters E, temp.cls stdeket)); OSMA_SWE_CALL(suddhimopy(clusters Einv, temp. yDericeTelebr1)); OSMA_WE_CALL(suddhimopy(clusters metheralis);

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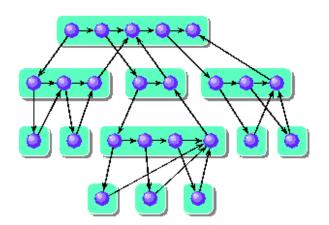
// Capy the likelihood totals from each block, sum them up to get a total DM_UME_CALL(outfourther/memory_like/)); DMB_DME_CALL(outfourther/memory_likelihood a likelihood a likelihood and likelihood a likeli

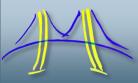


- Python AHC code is within 1.25x of pure C/CUDA implementation performance
- C/CUDA AHC (from winter retreat) 250x realtime
- SEJITized AHC ~ 200x realtime
- Time lost in:
 - Outer loop and GMM creation in Python
 - Data copying overhead from CPU to GPU
 - GMM scoring in Python

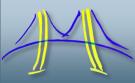


- We have implemented the Cilk backend for GMM training
- ASP selects version based on available hardware
- Current implementation ~100x realtime
- 5-10% C code reused
- All specializer infrastructure reused

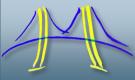




- Specializes to two types of platforms (multi-core CPU, Nvidia GPU) to support portability
 - Exact same application code
- Reuse of infrastructure:
 - Specializer creation and code variant selection mechanism reused
- Maintaining the code for next generation of hardware
 - Task of specializer writer, transparent to the application developer

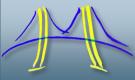


- SEJITized GMM training in Speaker Diarization component of Meeting Diarist
- Specialized covariance matrix computation with code variant selection to two platforms
- Currently a factor of 1.25x slower than pure C/CUDA implementation (200 x faster than realtime)
- Future work:
 - Further specialize train kernel
 - SEJITize other components
 - Improve code variant selection mechanism

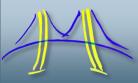


Thank you!

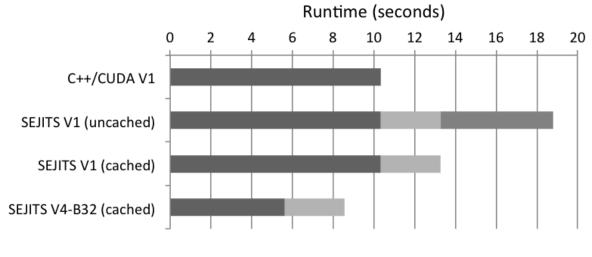
Questions?

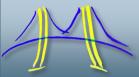


Backup Slides



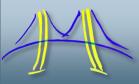
- Initial invocation 81% overhead dude to complier invocations
- Future runs using automatically determined optimal code variant achieve 17% performance *improvement* over the original GPU implementation (V1)





ASP vs Auto-tuning Libraries

	ATLAS	FFTW, Spiral, OSKI	ASP/GM M	ASP/Sten cil	Delite/ OptiML	Copperhe ad
Autuning of code						
Based on runtime information						
Based on higher- order func						
Using reusable framework						
Embedded in HLL						

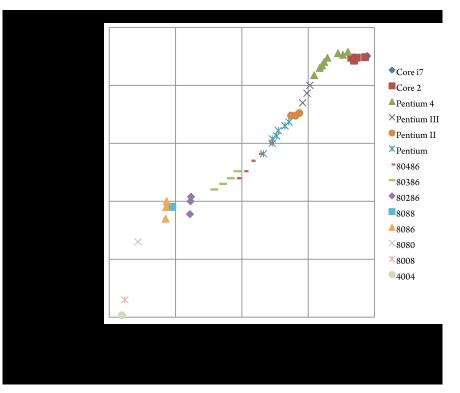


The shift to parallel processing

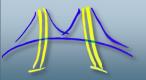
Parallel processing is here

** This shift toward increasing parallelism is not a triumphant stride forward based on breakthroughs in novel software and architectures for parallelism; instead, this plunge into parallelism is actually a retreat from even greater challenges that thwart efficient silicon implementation of traditional uniprocessor architectures.

- The Berkeley View

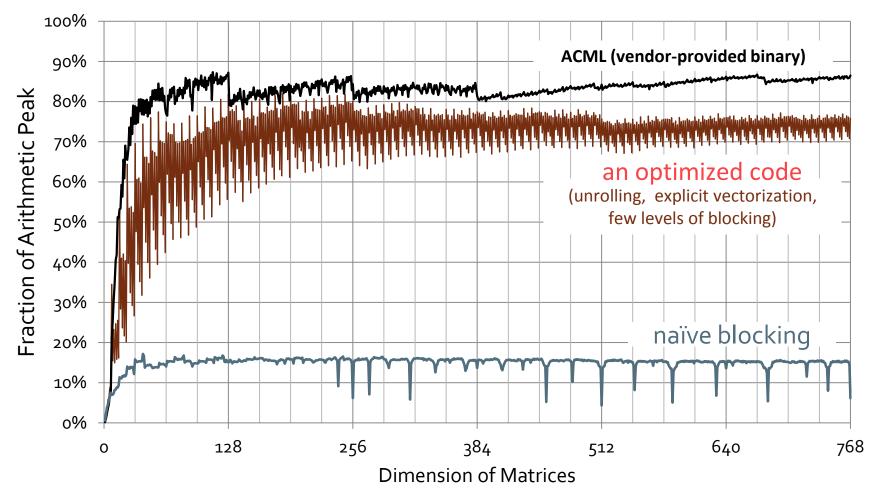


Intel Processor Clock Speed

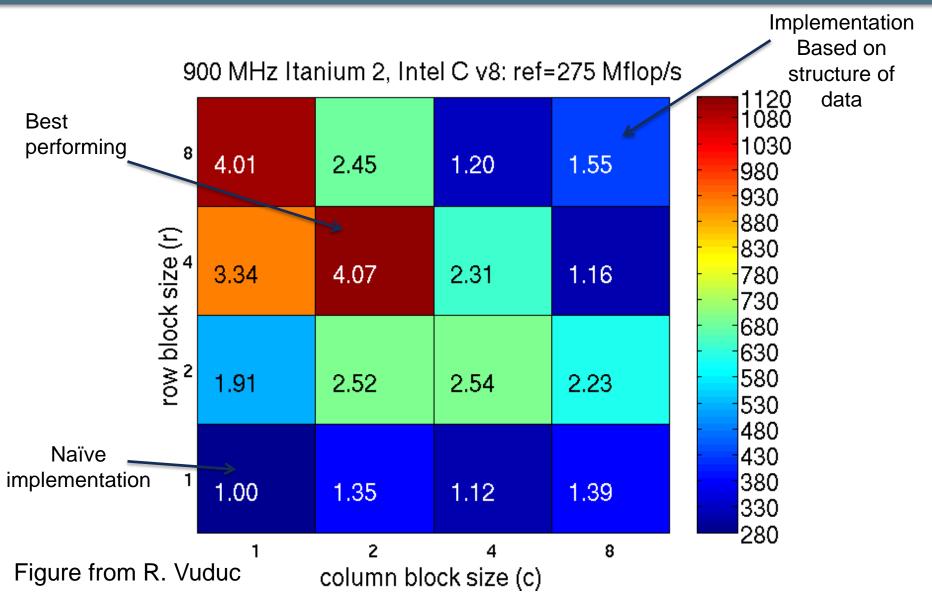


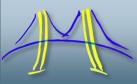
Writing Fast Code is Hard



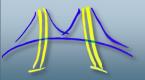


Finding Best Implementation is Hard

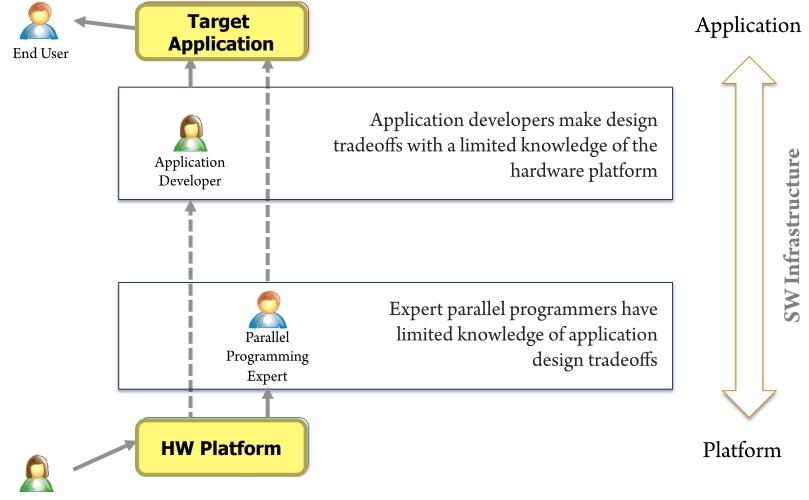




- Scientists and domain experts prefer to use highlevel languages such as Python or MATLAB
- However, to achieve sufficient performance, computationally-intensive parts of applications must eventually be rewritten in low-level languages
- In addition, parallel platform details and input parameters determine the best-performing parallel implementation



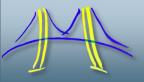
Implementation Gap



Hardware Architect

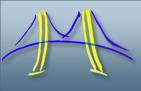


- SEJITS approach
- Gaussian Mixture Model & Applications
- Covariance Matrix Computation & Code Variants
- Specialization
- Results
- Conclusion & Future Work



Key Idea: Generate, compile, and execute high performance parallel code at runtime using code transformation, introspection, variant selection and other features of high-level languages.

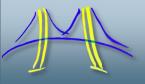
Invisibly to the user.



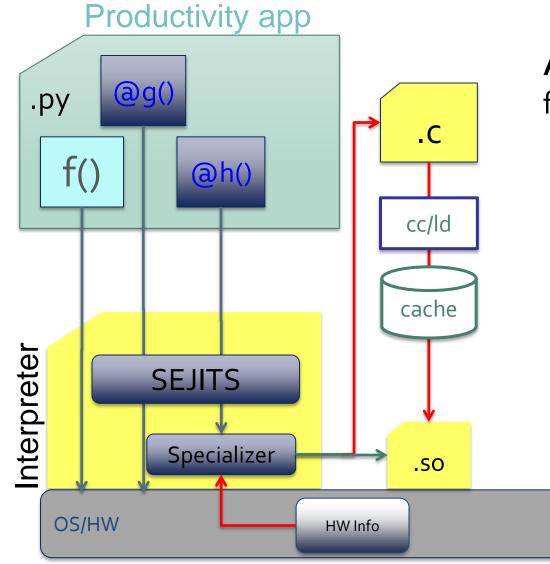
Selective Embedded JIT Specialization (SEJITS)

- Leverage patterns to bridge productivity and efficiency
- PLL (productivity-level language, eg Python) for applications
- "Specializers" generate ELL (efficiency-level language) code targeted to hardware
 - Code generation can happen at runtime
 - Specializers can incorporate autotuning
 - Think: pattern-specific embedded DSLs
- ELL performance with PLL effort

ΛΛ



Selective Embedded JIT Specialization (SEJITS)

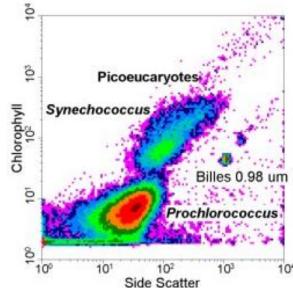


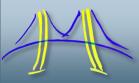
ASP – A SEJITS for Python

Applications of Gaussian Mixture Models

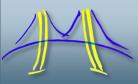
Applications

- Can be used to cluster/classify any sequence of observations
- Speech Recognition speaker classification, acoustic modeling for speech recognition
- Computer Vision image segmentation, hand writing recognition
- Biology flow cytometry
- Data mining topic classification in web documents
- Many more...

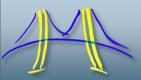




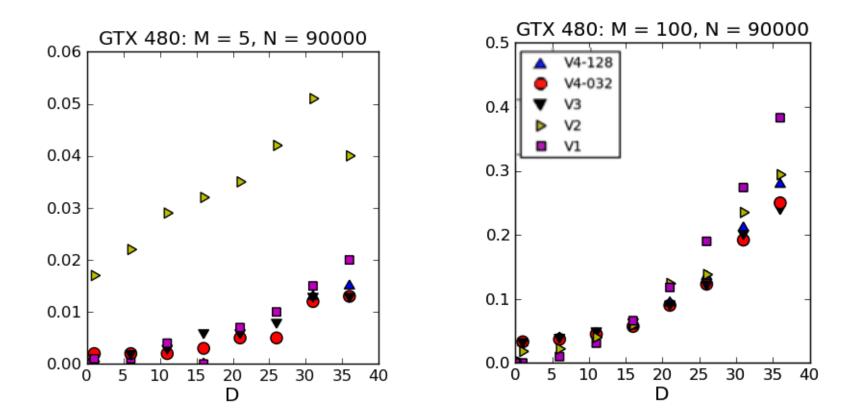
- Application example Agglomerative Hierarchical Clustering for Speaker Diarization
 - Uses GMMs to represent distribution of audio features for speakers in a recorded meeting
 - Iteratively trains GMMs using different number of components each time and measuring which number of components best fits the data
 - Number of components in the best GMM corresponds to number of speakers in the meeting



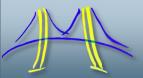
- ASP framework encapsulates code variant selection mechanisms and handcrafted templates to:
 - Allow domain expert to stay in the high-level language domain and focus on the application
 - Obtain high performance from expert-tuned code
- Example in Gaussian Mixture Model Applications
- Performance benefit of specialization outweighs the overhead of Python and the JIT process
- Expand to:
 - more platforms, applications, patterns
 - other code variant selection mechanisms



GTX₄80 – Varying D

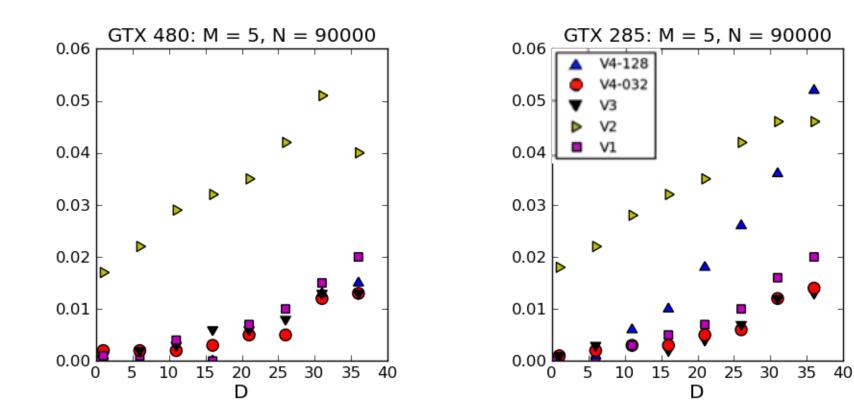


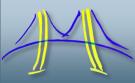
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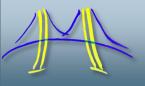
Results – Version Comparison (Raw CUDA)

GTX285 vs. 480





- ASP framework
 - C and CUDA compiling with CodePy (using PyCuda)
 - PyUBLAS to eliminate copies between C and Python
 - Version selection based on previous timings
- Evaluation platforms:
 - GTX480 (Fermi)
 - I4 SM, 32 SIMD, 48K shared mem, 3GB DRAM
 - GTX 285
 - 30 SM, 8 SIMD, 16K shared mem, 1GB DRAM
 - CUDA SDK 3.2
 - NVCC 3.2



Covariance Matrix Computation – Code Variants

