Performance Tuning of Deep Learning Framework Chainer on the K computer

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Introduction

Recently GPUs has become a popular platform for executing deep learning (DL) workloads. We revisit the idea of doing DL on CPUs, especially massively parallel CPU clusters (supercomputers). In anticipation of deployment of the Supercomputer Fugaku with much more DL capable CPUs, we investigate which optimizations can be already done using the K computer, current leadership computing facility and predecessor to the Supercomputer Fugaku. We use Chainer as a deep learning framework of choice.



- Profiling results of Chainer using cProfile+gprof2dot are [Fig.1][Table.1].
 - Total execution time of 10.311 s breaks down as follow:
 - Adam optimizer [adam.py]: 84%1.
 - numpy.dot called from linear.py: 11%2
 - Other parts (ReLU and Python control and so on.): 5%.

Adam optimizer

Optimizer consumed 84% of execution time and ran with 0.04% efficiency as measured using Fujitsu hardware counters.

tottime

8,405.11

ncalls

72000

28000

218000

24000

7102000 1,122.61

- The dominant operation is square root of matrix elements called from NumPy for filter update.
- This function in NumPy was implemented with automatic C language code generation, and thus difficult to tune directly...
- We rewrote all calculations in Adam using vectorized Fortran library and SIMD conversion and software pipelining (SWPL).
- In filter update calculations many values happened to be close to zero(denormalized number), raising underflow exceptions. We recompiled Python with an option forcing to truncate such numbers to zero.

Table.2 numPy.dot calculation.

gemm size(M,N,K)

We have also applied SWPL and masked SIMD using Fortran Library to implement ReLU activation function.

numpy.dot GEMM convolution

- achieves only 7.76% peak performance.
- NumPy was compiled against vectorized (using SSI II) but single-threaded Fujitsu BLAS library.
 - We modified numpy.dot to call multi-threaded version.

104	7200 4482481.0	622.6	21.2 m += (1 - hp.beta1) * (grad - m)
105	7200 4366618.0	606.5	20.6 v += (1 - hp.beta2) * (grad * grad - v)
	7200 7337882.0	1019.2	34.6 param.data -= hp.eta * (self.lr * m / (numpy.sqrt(vhat) + hp.eps)
113	7200 4864633.0	675.6	23.0 hp.weight decay rate * param.data)

Fig.2 Line costs of Adam optimizer region.

Table.1 ASIS Chainer Profiler (cProfile) result on the K computer.

0.117 optimizers/adam.py:91(update_core_cpu)

0.011 (method 'dot' of 'numpy.ndarray' objects)

elapse [s] efficency %

6.73% 8.90%

429.3[s]

283.2[s]

380.6[s]

0.007 function node.py:201(apply)

· Original Chainer Ver.4.4.0 profile for MNIST sample (unit=1,000, epoch=20) on the K compute

percall

percall cumtime

0.002

0.000

0.001

8,421.99

1,421.38

47.38

32.05

		linear.py:33(forward)	y=x.dot(W.T)	(100, 1000, 784), (100, 1000, 1000), (100, 10, 1000)	x42000
		linear.py:96(forward)	gx=gy.dot(W)	(100, 1000, 10), (100, 1000, 1000)	x24000
4.	other cost	linear.py:145(forward)	gW=gy.T.dot(x)	(10, 1000, 100), (1000, 1000, 100), (1000, 784, 100)	x36000
_	The ratio of the other	costs such as Rel II	calculations a	nd Python control regions in ASIS Chainer is	small (4

calculation

- ainer is small (4.5%).
- These regions almost not vectorized (SIMD) and parallelized.

We adopted **FlatMPI** on these regions for threads parallelization. Table.3 Elapsed time of tuning stage.

name	tuning	GEMM	Adam	other	total
origi	inal	1,189.1	8,657.4	465.2	10,311.7
underflow	Kfast(Kns)	990.2	1,681.9	372.2	3,044.4
GEMM	thread	236.5	8,386.9	336.9	8,960.2
Adam	FortranLib.	1,052.6	780.7	439.6	2,272.9
al	l	194.9	41.3	399.0	635.2
FlatMPI (8proc.)	131.9	40.6	78.6	283.4

Table. 4 Performance efficiency of GEMM and sqrt.

original 7.76% 0.04% Inderflow Kfast(Kns) 9.31% 0.19% GEMM thread 38.81% 0.48% Adam FortranLib. 8.76% 0.92% □ 47.03% 17.89% FlatMPI (8proc.) 70.05% 18.64%
GEMM thread 38.81% 0.48% Adam FortranLib. 8.76% 0.92% all 47.03% 17.89%
Adam FortranLib. 8.76% 0.92% all 47.03% 17.89%
all 47.03% 17.89%
FlatMPI (8proc.) 70.05% 18.64%

Table.5 Profiler result after all tuning

· Tunea Cr	· luned Chainer ver.4.4.0 profile for MN1S1 sample (unit=1,000, epoch=20) on the K computer					
ncalls	tottime	percall	cumtime	percall	filename:lineno(function)	
102000	166.80	0.002	166.80	0.002	{method 'dot' of 'numpy.ndarray' objects}	
72000	60.73	0.001	104.47	0.001	optimizers/adam.py:53(adam_kro2)	
218000	38.49	0.000	440.19	0.002	function_node.py:201(apply)	
12000	18.17	0.002	292.06	0.024	variable.py:968(_backward_main)	
364000	16.73	0.000	21.16	0.000	numpy/ctypeslib.py:196(from param)	

Speedups by each tuning step [Fig.3]:

- SSL II thread parallel BLAS effect in numpy.dot: 1.15x
- SWPL effect by Fortran library for Adam's sqrt: 4.54x
- floating point underflow control effect: 3.38x
- FlatMPI effect: 2.24x
- Total speedup: 36.4x

Fig.1 Call graph sample of Chainer

on the K computer

Result

- Efficiency improvements [Table 4]:
- Performance efficiency of the numpy.dot: 7.76% \rightarrow 38.81% \rightarrow 47.03% \rightarrow 70.05% (9.03x)
- Performance efficiency of the Adam optimizer: 0.04% → 0.92% → 17.89% → 18.64% (511x)

Performance tuning of Chainer Cost distribution of Chainer elapse 9 elapsed[12,000.0 on the K computer on the K computer 100% 90% **⊞**MPI 10,000.0 80% GEMN GEMM 70% 60% 8,657.4 40.6 6,000.0 4,000.0 10/10 2.000.0 1,681.9 10% 0.0 0% underflow GEMM original

Fig.3 Elapsed time on the all tuning step

Fig.4 The change of cost distribution by the tuning effect.

Conclusion

There are some limitations on the use of Chainer on the K computer. It is necessary to prepare the learning data beforehand and to stage-in the data to an appropriate storage system. Moreover, since Python is in the shared storage, it takes time to load the library. However, it was confirmed that we can use the K computer for deep learning sufficiently as well as GPU. Now, we are preparing for the use of deep learning calculations on the Fugaku.

Parallelization by the ChainerMN

We also tried to evaluate ChainerMN-1.3.0 and released it to the K computer users as an R-CCS software.

- Scalability of ChainerMN on the K computer is [Fig.5].
 - Measurement conditions: MNIST sample (unit=1000, epoch=20 → 1epoch=600iter).
- Although it can be measured even using 600 proc. or more, we must take care of recurrence and consistency of results. It scales well up to about 200 processes for this MNIST sample
- It became possible to calculate in 10 seconds for MNIST sample on the K computer.

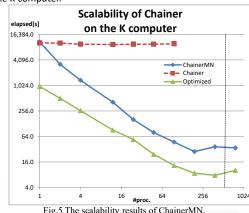


Fig.5 The scalability results of ChainerMN.